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A Small Molecule PAI-1 Functional Inhibitor Attenuates Neointimal Hyperplasia and Vascular Smooth Muscle Cell Survival by Promoting PAI-1 Cleavage

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Abstract

Plasminogen activator inhibitor-1 (PAI-1), the primary inhibitor of urokinase-and tissue-type plasminogen activators (uPA and tPA), is an injury-response gene implicated in the development of tissue fibrosis and cardiovascular disease. PAI-1 mRNA and protein levels were elevated in the balloon catheter-injured carotid and in the vascular smooth muscle cell (VSMC)-enriched neointima of ligated arteries. PAI-1/uPA complex formation and PAI-1 antiproteolytic activity can be inhibited, via proteolytic cleavage, by the small molecule antagonist tiplaxtinin which effectively increased the VSMC apoptotic index in vitro and attenuated carotid artery neointimal formation in vivo. In contrast to the active full-length serine protease inhibitor (SERPIN), elastasecleaved PAI-1 (similar to tiplaxtinin) also promoted VSMC apoptosis in vitro and similarly reduced neointimal formation in vivo. The mechanism through which cleaved PAI-1 (CL-PAI-1) stimulates apoptosis appears to involve the TNF-a family member TWEAK (TNF-a weak inducer of apoptosis) and it's cognate receptor, fibroblast growth factor (FGF)-inducible 14 (FN14). CL-PAI-1 sensitizes cells to TWEAK-stimulated apoptosis while full-length PAI-1 did not, presumably due to its ability to down-regulate FN14 in a low density lipoprotein receptor-related protein 1 (LRP1)-dependent mechanism. It appears that prolonged exposure of VSMCs to CL-PAI-1 induces apoptosis by augmenting TWEAK/FN14 pro-apoptotic signaling. This work identifies a critical, anti-stenotic, role for a functionally-inactive (at least with regard to its

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None declared.

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protease inhibitory function) cleaved SERPIN. Therapies that promote the conversion of fulllength to cleaved PAI-1 may have translational implications.

Keywords

Apoptosis; carotid stenosis; SERPIN; vascular injury; PAI-1

1. Introduction

Vascular restenosis, or blood vessel re-narrowing after percutaneous coronary intervention, is a combinatorial pathological consequence of increased vascular smooth muscle cell (VSMC) migration and proliferation combined with decreased apoptosis [1,2]. Few treatment options for vascular restenosis exist aside from additional catheterization. Among several factors implicated in the vascular response to injury, plasminogen activator inhibitor-1 (PAI-1), a member of the serine protease inhibitor (SERPIN) superfamily and the major physiologic regulator of the plasmin-based pericellular proteolytic cascade, is perhaps the most prominent.

PAI-1 exists in three distinct conformations [3,4]. While initially synthesized as an active SERPIN, PAI-1 spontaneously converts to a latent form (active half-life ~2 h at 37°C, pH 7.4) which is unable to inhibit uPA or tPA catalysis. This rather short half-life can be extended 2-to-10 fold, however, upon binding of PAI-1 to the somatomedin B domain of vitronectin where it also impacts integrin-matrix interactions and downstream signaling pathways [5–8]. During engagement of PAI-1 with its target proteases, the sissile bond in the reactive center loop (RCL) is cleaved to form a covalent ester bond between a serine hydroxyl group of the enzyme and a PAI-1 carboxyl group. Upon cleavage, the RCL N-terminus inserts into β -sheet A, while the C-terminus of the RCL forms strand s1C in β -sheet C producing a 70Å separation of the P1 and P1' residues, thereby deforming the complexed protease and rendering it inactive. A substrate form of PAI-1 exists as well in which PAI-1 is cleaved by its target proteases without formation of a covalent PAI-1:protease complex [3,9–11].

Due to the complexity of PAI-1 structure/function, several low-molecular weight antagonists of PAI-1 were developed to evaluate specific contributions of this SERPIN to disease pathology (12). Tiplaxtinin (PAI-039), a well-studied PAI-1 small-molecule inhibitor, attenuates asthmatic episodes, reduces hyperlipidemia and hyperglycemia and suppresses tumor angiogenesis [12–19]. The mechanism by which tiplaxtinin antagonizes the anti-fibrinolytic activity of PAI-1 appears to involve promotion of a substrate-like conformation resulting in PAI-1 cleavage and impaired uPA and tPA inhibition [20,21].

This paper reports that PAI-1 levels are elevated in injured VSMCs *in vivo*. Tiplaxtinin, which promotes PAI-1 cleavage, and elastase-cleaved PAI-1 (CL-PAI-1) both attenuate neointima formation in response to carotid artery ligation and stimulate plasmin-dependent, terminal VSMC apoptosis. CL-PAI-1, but not the functionally-stable full-length recombinant PAI-1 mutant 14-1b (FL-PAI-1), sensitizes VSMCs to TWEAK (TNF- α weak inducer of apoptosis)-induced apoptosis, a likely consequence of full-length PAI-1/LRP1-

mediated down-regulation of FN14 receptor expression. This work identifies a novel, antistenotic, role for cleaved PAI-1 in the tissue response to arterial injury. Therapies that promote the conversion of full-length to cleaved PAI-1 may have translational implications in the therapy of cardiovascular pathologies.

2. Experimental Procedures

2.1 Cell culture

Newborn rat arterial smooth muscle cells (RASMCs; gift from Dr. Peter A. Jones) were cultured in DMEM with 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin in a humidified 5% CO_2 atmosphere at 37°C. Human carotid artery SMCs (HuCASMCs; Cell Applications Inc., San Diego, CA) were grown in Smooth Muscle Cell Growth Medium containing 5% FBS, 1% penicillin/streptomycin and 0.5% each hEGF, insulin and bFGF/ heparin, conditions which maintain VSMC in the de-differentiated, synthetic phenotype.

2.2 Recombinant proteins

Endotoxin levels in full-length, stabilized recombinant 14-1b PAI-1 (FL-PAI-1), R76E-PAI-1 (LRP1-binding deficient mutant; Molecular Innovations, Novi, MI), human neutrophil elastase (Abcam, Cambridge, UK) and high-molecular weight two chain uPA (Seiksui Diagnostics, LLC, Lexington, MA) preparations were measured using the Limulus Amebocyte Lysate kit QCL-1000 (Lonza, Basel, Switzerland) and all found to be <0.14 EU/ml, the acceptable threshold [22]. Recombinant Human TWEAK/TNFSF12 was obtained from R&D Systems (Minneapolis, MN).

2.3 RNA interference

siRNA designed against human PAI-1 mRNA (5'-AAGGATGAGATCAGCACCACA-3') and a scrambled control (sc) siRNA (5'-AATTCTCCGAACGTGTCACGT-3') [23] were obtained from QIAGEN Inc. (Valencia, CA). HuCASMCs were transfected with 20 nM of PAI-1 siRNA or sc siRNA using Lipofectamine 2000 (Invitrogen).

2.4 Balloon Catheter Denudation and Carotid Artery Ligation

All animal protocols were approved by the Institutional Animal Care and Use Committee of Albany Medical College (IACUC approved protocol #13-09001) and conducted in accord with the EU Directive 2010/63/EU for animal experimentation. Rodents were housed in the College Animal Resource Facility, licensed by the USDA and New York State Department of Health, Division of Laboratories and Research. Male Sprague-Dawley rats (400~450 g, Taconic Farms, Germantown, NY) were anesthetized by i.p. injection of ketamine (0.1 mg/gm) and xylazine (0.01 mg/gm), the left common carotid artery exposed by a midline cervical incision and blunt dissection performed alongside the artery with dull forceps to expose the carotid bifurcation into the internal/external branches. After blood flow cessation by arterial clamping, a 2F Fogarty balloon catheter (Edwards) was introduced *via* a small arteriotomy in the external carotid and advanced to the common carotid artery [24]. The balloon was inflated by 1.6 atm pressure, inserted and withdrawn three times. For carotid ligation, FVB/NJ mice (Jackson Labs) were anesthetized by i.p. injection of ketamine (0.1 mg/gm) and xylazine (0.01 mg/gm). Following site preparation, the left carotid artery was

exposed with a small 8–10 mm midline incision in the neck and blunt dissected to free the left common carotid and branches from surrounding tissue. The common carotid artery was ligated just proximal to the internal and external carotid bifurcation with a 6-0 sterile silk suture. Animals were treated with tiplaxtinin (3 mg/kg; oral gavage), full-length (FL-PAI-1; 3 mg/kg; i.p. injection), and cleaved PAI-1 (CL-PAI-1; 3 mg/kg; i.p. injection), or vehicle control once daily for 14 days following ligation.

2.5 Northern Blotting

Total RNA (10 µg) from the left (denuded) and (right) uninjured carotid arteries was separated on 1.2% agarose/formaldehyde gels, transferred to Nytran membranes by capillary action in 10x SSC (3 M NaCl, 0.3 M Na citrate, pH 7.0) and immobilized by UV crosslinking prior to incubation for 2 h at 42°C in prehybridization buffer (50% formamide, 5X Denhardt's reagent, 1% SDS, 100 µg/ml sheared/heat-denatured salmon sperm DNA [ssDNA], 5X SSC). RNA blots were incubated with random-primed ³²P-dCTP-labeled cDNA probes (5×10^6 cpm) to rat PAI-1 or A50 (loading control) for 16 h at 42°C in hybridization buffer (50% formamide, 2.5x Denhardt's reagent, 1% SDS, 100 µg/ml ssDNA, 5x SSC, and 10% dextran sulfate). Membranes were washed in 0.1X SSC/0.1% SDS at 42°C, followed by three 15 min washes in 0.1 x SSC/0.1% SDS at 55°C and exposed to X-OMAT AR-5 film using intensifying screens. Probe was removed from blots by washing in 55% formamide, 2X SSC, 1% SDS for 1 h at 65°C and rinsed in 0.1 x SSC/0.1% SDS prior to rehybridization with A50.

2.6 Immunohistochemistry

Paraffin-embedded tissue sections (5 μm) were stained with hematoxylin and eosin (H&E). For immunohistochemistry, sections were incubated with primary antibodies to smooth muscle α-actin (Sigma-Aldrich, St. Louis, MO; 1:6,400 dilution) and PAI-1 (Affinity Bioreagents, Waltham, MA; 1:1,000 dilution), washed, incubated with biotinylated antirabbit IgG and avidin D horseradish peroxidase (1:1,000 dilution) and color reactions developed using an ABC kit prior to mounting coverslips in Vectashield medium (Vector Labs, Burlingame, CA). Isotype IgG controls were included. Sections were analyzed using Nanozoomer 2.0RS digital microscope equipped with NDP 2.2.1 software (Hamamatsu Photonics, Hamamatsu, Japan).

2.7 Flow Cytometry

Fluorescence intensity was measured using a FACS LSRII (BD Biosciences, San Jose, CA). Ten thousand events were counted for each sample and experiments analyzed using FlowJo software (Tree Star Inc., Ashland, OR).

Cell Cycle Analysis—After treatment, RASMCs and HuCASMCs were harvested by trypsinization, centrifugation, washed twice with ice-cold phosphate-buffered saline (PBS) and fixed in 70% ethanol for 1 h. Fixed cells were washed twice with ice-cold PBS and incubated for 2 h in the dark in PBS containing 0.1% Triton-X100, 20 µg/ml Propidium Iodide (PI; Sigma Aldrich, St. Louis, MO) and 10 µg/ml RNase A.

Annexin-V—RASMCs, treated as indicated in the text, were trypsinized, collected and stained with FITC-conjugated Annexin-V (Calbiochem, Darmstadt, Germany).

Cleaved Caspase-3—Treated RASMCs were harvested by trypsinization, collected by centrifugation, washed with 0.2% BSA in PBS and fixed with 4% paraformaldehyde for 15 min. Cells were permeabilized in 90% cold methanol for 30 min, followed by two washes and incubated with antibodies to cleaved caspase-3 (1:200 dilution, Asp175; Cell Signaling, a, MA) followed by Alexa-Fluor 488-tagged IgG (1:1000 dilution) for 60 min each. Cells were washed prior to fluorescence measurements as described above.

FN14 surface expression—After treatment, RASMCs and HuCASMCs were harvested by trypsinization, collected and incubated 10% FBS/phosphate-buffered saline (PBS) for 15 min. Cells were pelleted and washed in 0.1% BSA/PBS, followed by incubation with FN14 antibody or mouse IgG (5 μg/ml in 0.1% BSA/PBS; Santa Cruz Biotechnology, Dallas, Texas) followed by Alexa-Fluor 488-tagged IgG (1:1000 dilution for 1 h each). Cells were washed, resuspended in 2% paraformaldehyde/PBS and fluorescence measurements assessed as described above.

2.8 PAI-1-GFP chimera expression construct

Human PAI-1 promoter sequences (-800 to +71) were PCR-amplified for 30 cycles using the p800-Luc reporter plasmid as a template and Platinum *Taq* polymerase. This fragment was gel-purified for subsequent cloning into the *SacI/KpnI* sites of the promoter-less GFP expression vector pEGFP-1 (Clontech, Palo Alto, CA). For preparation of a PAI-1-GFP chimeric expression construct, the full-length human PAI-1 coding sequence (approximately 1.3 kb) was derived by RT-PCR from total RNA isolated from human foreskin fibroblasts, amplified by PCR, gel-purified and expressed as a GFP fusion protein by T4 ligase insertion into the *Bam*HI/*AgeI* site of the PAI-1 promoter-derived pEGFP-1 vector. All constructs were sequence verified. RASMCs were seeded into 35 mm dishes and allowed to reach a density of 1×10^5 cells/cm² prior to transfection with $1-2 \mu g$ DNA using Lipofectamine-Plus.

2.9 Apoptosis Assay

Annexin-V—RASMCs cultured in 96-well plates were treated with the indicated concentrations of tiplaxtinin in complete media for 24 h. Cells were stained with FITC-conjugated Annexin-V and apoptosis measured using a Biotek Synergy 2 fluorescent microplate reader and Biotek Gen5 software. The vehicle control (DMSO) did not induce apoptosis.

Caspase-3 activity assay—Asynchronously growing HuCASMCs were treated as indicated in complete media for 6 h. Cells were harvested, lysed, and centrifuged to remove cellular debris. Aminomethylcoumarin (AMC)-derived substrate Z-DEVD-AMC was added to cell lysates and incubated for 30 minutes according to manufacturer's instructions (EnzChek® Caspase-3 Assay Kit #1, Life Technologies, Carlsbad, CA). Fluorescent emission (excitation/emission maxima ~342/441 nm) was read every hour using a Biotek Synergy 2 fluorescent microplate reader and Biotek Gen5 software.

Live/Dead assay—Asynchronously growing HuCASMCs were treated as indicated in complete media for 24 hrs. Cell-permeant calcein AM which produces green fluorescence (ex/em ~495/~515 nm) in live cells and ethidium homodimer which only enters damaged membranes and produces a red fluorescence (ex/em ~495 nm/~635 nm; Live/Dead Viability/Cytotoxicity assay for mammalian cells, Life Technologies, Carlsbad, CA) were added directly in complete media at final concentrations of 2 μ M and 4 μ M, respectively. Cells were incubated with the Live/Dead reagents for 30 minutes, then counted. Cells positive for ethidium homodimer were scored as apoptotic and normalized against calcein AM (total cells).

2.10 Generation of Elastase-Cleaved PAI-1

Full-length (FL-PAI-1,14-1b stable PAI-1 mutant; Molecular Innovations, Novi, MI) was incubated with a 0.1 molar equivalent of human neutrophil elastase (Abcam, Cambridge, UK) in 100 mM Tris-HCl, pH 7.5, buffer containing 500 mM NaCl as described previously [3] to generate cleaved PAI-1 (CL-PAI-1). For each reaction time point, an aliquot was withdrawn and treated with 10 mM PMSF prior to SDS-PAGE and silver staining to confirm cleavage. Reaction products, without added PMSF, were stored at -80°C. CL-PAI-1 was also generated by incubation with Sepharose bead-immobilized porcine pancreatic elastase (Molecular Innovations, Novi, MI).

2.11 Tiplaxtinin:PAI-1:uPA Interactions

FL-PAI-1 (500 nmol/L) was incubated with tiplaxtinin or vehicle alone in 10 mM HEPES, pH=7.4, 150 mM NaCl and 0.5% Tween-20 buffer for 15 min before addition of 500 nM high-molecular weight uPA for a 5 min incubation at 37°C [20]. Enzymatic activity was stopped using 10 mM (final concentration) PMSF and reaction products separated by SDS-PAGE.

2.12 Western Analysis

Cells were washed with PBS and lysates prepared in 60 mM Tris HCl, pH=6.8, and 2% SDS. Protein concentration was determined using BCA Assay (Pierce Scientific, Rockford, IL) and 40 µg protein loaded for each sample. Western blotting was as described [4]. Blot analysis used the ChemiDoc MP System (Bio-Rad, Hercules, CA). Antibodies were: caspase-8 (1C12), caspase-3, pAkt S473 (D9E), pan Akt (11E7), PARP (from Cell Signaling, Beverly, MA), plasminogen (#364R, Seiksui Diagnostics, LLC, Lexington, MA) and FN14 (ITEM-4), TWEAK (S-20), GAPDH, ERK2 (C-14), actin (I-19) (Santa Cruz, Dallas, TX).

2.13 Statistics

Values are represented as mean \pm standard error of the mean (SEM). Student's T-tests were used to determine significance for experiments containing two experimental conditions. One-way ANOVAs with a Tukey's post-hoc analysis were used for analysis of experiments containing three or more conditions. Experiments that monitored dependent variables over time were analyzed using repeated-measures ANOVA. P-values <0.05 were considered statistically significant.

3. Results

3.1 PAI-1 expression is induced in vessel injury in vivo

To examine the relationship between PAI-1 expression and the pathogenesis of neointimal hyperplasia, PAI-1 mRNA levels were assessed in rats in which stenosis was induced in response to balloon catheterization. PAI-1 transcripts were markedly elevated in the catheter-denuded carotid artery as compared to the contralateral control vessel (Fig. 1A). The well-accepted carotid ligation model of arterial injury was used to establish the specific topographic localization of PAI-1 in stenotic lesions. The PAI-1-positive neointima region co-localized with smooth muscle cell α -actin-expressing cells (Fig. 1B) suggesting that the pathological response to vascular injury *in vivo* is associated with increased PAI-1 in the expanded VSMC cohort.

3.2 Tiplaxtinin initiates a two-step inhibition of PAI-1 antiproteolysis and stimulation of VSMC apoptosis

Tipaxtinin, a small molecule inhibitor of PAI-1 function, blocks PAI-1 anti-proteolytic activity in a two-step process. In an *in vitro* cell-free analysis, tiplaxtinin interferes with uPA::PAI-1 complex formation (Fig. 2A, **upper bands**) while stimulating (presumably a substrate-like) PAI-1 cleavage (Fig. 2A, **middle bands**). Compared to control untreated cells in which only one PAI-1 band was detected at 47 kDa, a low molecular weight (43 kDa) PAI-1 species was also evident indicating that tiplaxtinin stimulated PAI-1 cleavage by cultured cells as well (Fig. 2B).

Tiplaxtinin-induced cleavage, which generates a non-functional SERPIN (at least with regard to its anti-proteolytic activity), may have significant physiologic consequences. Indeed, PAI-1 is a pro-survival factor in several cell types and both engineered overexpression of wild-type full-length PAI-1 (Fig. 3A,B) and exogenous addition of a stable recombinant PAI-1 mutant (14-1b; FL-PAI-1) (Fig. 3C,D) increased pAKT levels in VSMCs. Since tiplaxtinin significantly blocked the pAKT response to FL-PAI-1 (Fig. 4A), and whereas CL-PAI-1 was markedly deficient (by 50% relative to FL-PAI-1) in pAKT inducibility, these data suggested that a catalytically intact SERPIN was required for Akt activation. It was important, therefore, to evaluate the effect of this small molecule inhibitor of PAI-1 function on VSMC survival. Annexin-V measurements indicated that tiplaxtinin dose-dependently induced RASMC apoptosis (Fig. 4B). Previous reports indicated, moreover, that the tiplaxtinin-binding site on PAI-1 approximates the vitronectin recognition site [20]. Despite the fact that the majority of circulating PAI-1 is bound to vitronectin, which prolongs the active half-life and stabilizes the bioavailability of this SERPIN, RASMCs plated on vitronectin-coated plates remained susceptible to tiplaxtinin-stimulated apoptosis (Fig. 4C).

3.3 Tiplaxtinin induces VSMC apoptosis by promoting a switch from full-length to cleaved PAI-1

Since tiplaxtinin stimulates PAI-1 cleavage as well as VSMC apoptosis (Fig. 4B), it was necessary to determine if cleaved PAI-1 mimicked the tiplaxtinin-induced apoptotic response. Advantage was taken of the time-dependent cleavage of PAI-1 by elastase, a well-

established model to evaluate the impact of cleaved PAI-1(CL-PAI-1) [25] (Fig. 4A, **compare shift in total PAI-1 [solid arrow] to cleaved PAI-1 [dotted arrow]**). Incubation of HuCASMC with CL-PAI-1 but not full-length (FL-PAI-1) resulted in a dramatic increase in apoptosis; the extent of the apoptotic response was dependent on the efficacy of PAI-1 cleavage (Fig. 5B). Identical results were obtained using Sepharose bead-immobilized elastase to generate PAI-1 cleavage (**not shown**). CL-PAI-1, moreover, significantly increased the level staurosporine-induced active caspase-3 (Fig. 5C). In contrast to FL-PAI-1-treated and control mice, i.p.-injected CL-PAI-1 and gavage-delivered tiplaxtinin (used separately) decreased neointimal thickening in response to carotid artery ligation (Fig. 5D–F). These data suggest that conversion of full-length to cleaved PAI-1 attenuates intimal hyperplasia by stimulating VSMC apoptosis.

3.4 Cleaved PAI-1 promotes plasmin-dependent VSMC apoptosis

As plasmin activation stimulates apoptosis [26], it was important to determine if CL-PAI-1 induces a plasmin-dependent apoptotic response. Addition of either tiplaxtinin or CL-PAI-1 increased the conversion of plasminogen to plasmin (Fig. 6A, *left*) to approximately the same level of plasmin activation as a result of PAI-1 knockdown (Fig. 6A, *right*). Poly (ADP-ribose) polymerase (PARP) cleavage, a down-stream effector of caspase-3, was also up-regulated in response to CL-PAI-1 while Tranexamic Acid, a potent inhibitor of plasmin generation, attenuated the apoptotic response suggesting that CL-PAI-1 stimulates apoptosis through plasmin-dependent caspase-3 activation (Fig. 6B). These data are consistent with additional findings that another plasmin inhibitor (aprotinin), a uPA inhibitor (amelioride), and the pan-MMP inhibitor (prinomastat) also reduced cleaved PAI-1-stimulated apoptosis (**not shown**). Plasminogen alone had no effect on cell viability; however, in the presence of tiplaxtinin, the increase in VSMC apoptosis was likely a consequence of decreased uPA inhibition and increased plasmin activation (Fig. 6C). These data confirm that CL-PAI-1 stimulates VSMC apoptosis in a plasmin-dependent manner.

3.5 Cleaved PAI-1 upregulates FN14 and TWEAK expression while Full-length PAI-1 downregulates FN14 surface expression via LRP1 and prevents TWEAK stimulated apoptosis

To determine the signaling intermediates involved in cleaved PAI-1-induced apoptosis, the activation of caspase-8 in response to CL-PAI-1 and tiplaxtinin (used separately) suggest the potential involvement of death receptors (Fig. 7A). One pathway recently identified to regulate PAI-1 in atherosclerotic lesions is the fibroblast growth factor-inducible 14 (FN14) receptor and its cognate ligand, TNF-α weak inducer of apoptosis (TWEAK) [47]. FN14 and soluble TWEAK levels were up-regulated in response to both CL-PAI-1 and tiplaxtinin while neither had any effect on total cell-associated TWEAK (Fig. 7B). Importantly, decreased circulating soluble TWEAK levels have been associated with poor prognostic outcomes in patients with cardiovascular and chronic kidney diseases [50] and based on these findings, it was necessary to determine if PAI-1 (cleaved or full-length) might modulate surface FN14 levels and signaling. HuCASMCs exhibited a significant down-regulation of FN14 in response to FL-PAI-1, but not CL-PAI-1 (Fig. 8A). PAI-1 recognizes the endocytic low-density lipoprotein receptor-related protein-1 (LRP1) in a multimeric complex that includes uPA/tPA and integrins as binding partners inducing their internalization and lysosomal degradation [28]. Since PAI-1-dependent activation of Akt

was inhibited by RAP (Fig. 8B) and LRP1^{-/-} cells were not inducible for PAI-1-initiated pAkt induction (Fig. 8C), it was necessary to determine if LRP1-binding activity was required for FN14 down-regulation. Unlike incubation of VSMCs with FL-PAI-1 which reduced FN14 immunoreactivity, addition of the R76E-PAI-1 mutant (which cannot bind LRP1 [29]) failed to down-regulate surface FN14 levels (Fig. 8D). HuCASMCs were treated with both TWEAK and either FL-PAI, CL-PAI-1, or R76E-PAI-1 to directly test if PAI-1 impacts TWEAK-mediated apoptosis. While TWEAK alone and in combination with FL-PAI-1 had no effect on apoptosis, both CL-PAI-1 and R76E-PAI-1 in addition to TWEAK significantly stimulated an apoptotic response (Fig. 8E). These data suggest that FL-PAI-1, but not CL-PAI-1, down-regulates FN14 via LRP1-mediated endocytosis preventing TWEAK-stimulated apoptosis.

4. Discussion

The relative abundance of the different conformational pools of PAI-1 may dictate whether VSMCs migrate or undergo apoptosis in response to injury. Indeed, in the context of cardiovascular disease, it appears that PAI-1 is both pro- and anti-restenotic [reviewed in 30] depending on the specific wound model, level of PAI-1 induction and vessel TGF- β 1 expression. Global PAI-1 deletion, over-reliance on application of only the full-length, active form of PAI-1 to assess vascular remodeling, and uncertainties as to the role of PAI-1 conformation-dependent processes, however, are major confounders.

It is increasingly evident that beyond its anti-proteolytic role, PAI-1 also functions as a multifunctional signaling "ligand" where it impacts cellular responses, including the migratory, proliferative and survival programs, at the site of injury [22,28]. The ability of active PAI-1 to inhibit apoptosis, however, is not due to its ability to bind uPA or signal through uPAR [31]. Consistent with the present findings that FL-PAI-1 activates Akt and that CL-PAI-1 induces VSMC apoptosis, a truncated PAI-1 (PAI-123) mutant, deleted in much of the heparin binding-domain and the RCL, also stimulated endothelial cell apoptosis [32,33]. It appears that an intact RCL is required for the pro-survival function of PAI-1. Indeed, application of CL-PAI-1 results in an apoptotic respose, both with and without additional death stimuli, while tiplaxtinin (10 µM), in the presence of FL-PAI-1 (40 nM), promotes apoptosis to a greater extent than tiplaxtinin alone. Since 20 µM tiplaxtinin generates the complete cleavage of 400 nM PAI-1 in 30 min, it is likely that under more prolonged exposure (24 h) tiplaxtinin effectively stimulates the cleavage of both exogenous recombinant and endogenous PAI-1 pools. Importantly, VSMCs show robust expression of tPA and uPA (not shown), a prerequisite for tiplaxtinin-mediated PAI-1 proteolytic cleavage.

Interestingly, elastase levels increase immediately after balloon-angioplasty, peak at one week and then decline [34] suggesting that lowered elastase activity levels, in the latter stages of vessel "repair" may contribute to a decrease in PAI-1 cleavage at the injury site resulting in reduced apoptosis and increased neointima expansion through both VSMC proliferation and migration. As elastase cleaves PAI-1 at the peptide bond between Val³⁴³- Ser³⁴⁴ (P4-P3) [25] and, whereas, tiplaxtinin promotes substrate behavior of PAI-1 allowing for a uPA/tPA protease attack at Arg³⁴⁶-Met³⁴⁷ (P1-P1') [35], cleavage at either of these

sites may be sufficient for promotion of apoptosis. Indeed, ligation-induced neointima formation was significantly attenuated by treatment with both tiplaxtinin and CL-PAI-1. Tiplaxtinin could be a useful therapeutic option in the context of elevated uPA and PAI-1 as it promotes a substrate-like conversion of PAI-1 [36–42].

Assessment of the mechanisms underlying the apoptotic response of VSMCs to CL-PAI-1 and tiplaxtinin implicated the extrinsic death receptor cascade since caspase-8 activation was a common event upon exposure to both. In the context of PAI-1 deficiency, the increase in active plasmin releases a pro-apoptotic soluble Fas ligand that stimulates endothelial cells apoptosis [43]. VSMCs, however, are relatively resistant to Fas ligand-stimulated death [44] and an antagonizing antibody to Fas ligand was unable to rescue VSMCs from tiplaxtinin or cleaved PAI-1-induced apoptosis (not shown). One candidate extrinsic apoptosis signaling pathway involved was the TWEAK/FN14 signaling axis since PAI-1 expression in atherosclerotic lesions appears responsive to exogenous TWEAK application [47]. TWEAK can activate its cognate receptor, FN14, either as a membrane-bound or solubilized ligand. The down-stream signaling and biological repercussions are tissue-type and contextdependent but have been implicated in apoptotic, proliferative and migratory processes [48].

The three forms of PAI-1 (full-length, latent and cleaved) interact with the endocytic low density lipoprotein receptor-related protein 1 (LRP1) and stimulate JAK/STAT1-mediated VSMC migration [22]. Unlike FL-PAI-1 which down-regulates surface FN14, a full-length PAI-1 mutant unable to bind LRP1 (R76E-PAI-1) [49], failed to reduce FN14 levels. Furthermore, addition of soluble, recombinant TWEAK sensitized VSMC to apoptosis in the presence of CL-PAI-1 and R76E-PAI-1, but not in cells exposed to FL-PAI-1. This is significant since, within the context of cardiovascular disease, it appears that circulating soluble TWEAK levels are negatively correlated with disease incidence [50] and tiplaxtinin and CL-PAI-1, but not FL-PAI-1, both up-regulate membrane-bound and soluble TWEAK levels. How CL-PAI-1 promotes TWEAK solubility is unknown. The protease furin, an inhibitory target of PAI-1, cleaves TWEAK; the involvement of other protease systems (e.g. plasmin) in TWEAK bioactivity are unknown, however it appears that CL-PAI-1 versus FL-PAI-1 differentially regulate several components of the FN14:TWEAK signaling axis to impact VSMC survival.

Collectively, these data are consistent with the hypothesis that PAI-1 regulates VSMC survival (Fig. 8). Full-length PAI-1 binds, and inhibits, uPA attenuating the conversion of plasminogen to plasmin while simultaneously down-regulating surface FN14 receptor expression via LRP1-mediated endocytosis. However, when the conformational pools of PAI-1 shift to increased levels of cleaved PAI-1 (either by tiplaxtinin or exogenous addition), conversion of plasminogen to plasmin is amplified. Both membrane-bound and soluble TWEAK expression is potentiated thereby stimulating FN14-TWEAK signaling leading to significant apoptosis. Since tiplaxtinin and CL-PAI-1 reduced neointimal lesion formation, therapies directed at manipulation of PAI-1 conformation may have applicability as a treatment option for restenosis

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7. Declaration of Funding

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Abbreviations

CL-PAI-1	cleaved PAI-1
FL-PAI-1	full-length PAI-1
HNE	human neutrophil elastase
HuCASMCs	human carotid artery smooth muscle cells
LRP1	low density lipoprotein receptor-related protein 1
pAkt	phospho-Akt
PAI-1	plasminogen activator inhibitor-1
PARP	poly (ADP-ribose) polymerase
Plg	plasminogen
RASMCs	rat arterial smooth muscle cells
RCL	reactive center loop
SERPIN	serine protease inhibitor
SERPINE1	serine protease inhibitor clade E member 1
ТРХ	tiplaxtinin
uPA	urokinase
VN	vitronectin
VSMCs	vascular smooth muscle cells

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Highlights

- PAI-1 expression is increased in the neointima of balloon-catheterized and ligated carotid arteries
- The small molecule PAI-1 inhibitor tiplaxtinin inhibits PAI-1/uPA complex formation and stimulates PAI-1 cleavage
- Tiplaxtinin and cleaved PAI-1, each independently, attenuate the neointimal response in the ligated carotid
- Tiplaxtinin and cleaved PAI-1 both stimulate VSMC apoptosis in vitro
- Prolonged exposure of VSMCs to cleaved PAI-1 induces apoptosis by augmenting TWEAK/FN14 pro-apoptotic signaling.

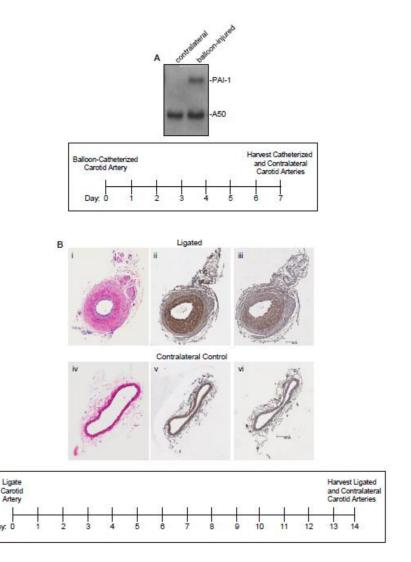


Figure 1. PAI-1 expression is elevated in smooth muscle cells of the neointimal compartment of injured vessels

(A) RNA was extracted from balloon catheter-denuded and contralateral control carotid arteries one week following injury. Northern blots were hybridized using ³²P-labeled cDNA probes to PAI-1 and A50 (loading control). (B) Paraffin sections (5 μ m) of ligated (i–iii) and contralateral control (iv–vi) carotids were stained with H&E (i, iv) or processed for immunohistochemical detection of smooth muscle cell α -actin (ii, v) or PAI-1 (iii, vi) 14 days after common carotid artery ligation. Scale bar=100 μ m.

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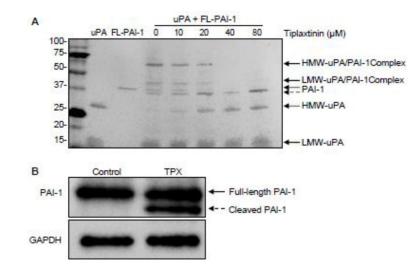


Figure 2. Tiplaxtinin inhibits PAI-1/uPA complex formation and promotes PAI-1 cleavage

(A) Human uPA was incubated with FL-PAI-1 and the indicated concentration of tiplaxtinin. Solid arrow indicates full-length PAI-1; dashed arrow denotes cleaved PAI-1.
(B) RASMCs were treated with tiplaxtinin or vehicle control in complete media for 24 h.
PAI-1 and GAPDH (loading control) in cell lysates were identified by western blotting.
Full-length and cleaved PAI-1 are highlighted by solid and dashed arrows, respectively.
TPX=tiplaxtinin.

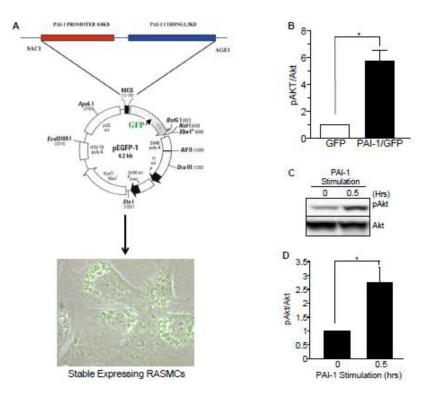


Figure 3. Vector-driven expression and exogenous addition of full-length PAI-1 stimulates Akt phosphorylation

(A) Schematic of a pEGFP-1-based vector in which a chimeric transcript consisting of 1.3 kb of PAI-1 coding sequences and GFP is expressed under the control of a 0.8 kb PAI-1 promoter. RASMCs were transfected with this PAI-1 coding sequence-GFP fusion construct under control of the PAI-1 promoter and stable expressing clones selected. (B) PAI-1 overexpression by 6- to 8-fold relative to GFP in RASMCs results in elevated pAkt levels.
(C) PAI-1 addition induces a rapid and transient Akt activation in RASMCs. At confluence, RASMCs were serum-starved for 48 hours followed by stimulation with full-length PAI-1 (40 nM) for 30 minutes and cellular lysates analyzed by western blot. (D) pAkt levels were normalized to total Akt levels; histogram illustrates mean ± SEM; asterisk=p<0.05; n=3.

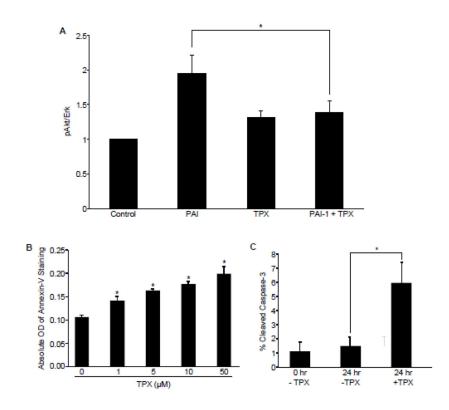


Figure 4. Tiplaxtinin inhibits PAI-1-induced Akt phosphorylation and promotes VSMC apoptosis

(A) RASMCs were serum-starved for 48 hours then treated with 40 nM PAI-1 with or without 10 nM tiplaxtinin. Cells were lysed, western blots probed for pAkt and normalized to Erk1. Data represent mean \pm SEM, N=3. Asterisk=p 0.05; n=3. (B) RASMCs, cultured in 96-well plates, were treated with various concentrations of tiplaxtinin in complete media for 24 h prior to incubation of FITC-conjugated Annexin-V antibodies; apoptosis was measured using a fluorescence microplate reader. Data are presented as mean \pm SEM; asterisks=p<0.05, n=3. (C) RASMCs were plated under reduced-serum conditions on tissue culture dishes pre-incubated with vitronectin and allowed to adhere and spread for 24 h. Cells were then either collected and fixed for immunocytochemical analysis of cleaved caspase-3 expression or treated with 10 μ M tiplaxtinin or vehicle control for 24 h prior to fixation. Cleaved caspase-3-positive cells were scored as apoptotic and normalized to total cells (as assessed using DAPI staining). Data plotted are the percent mean \pm SEM; asterisk=p<0.05, n=3. TPX=tiplaxtinin.

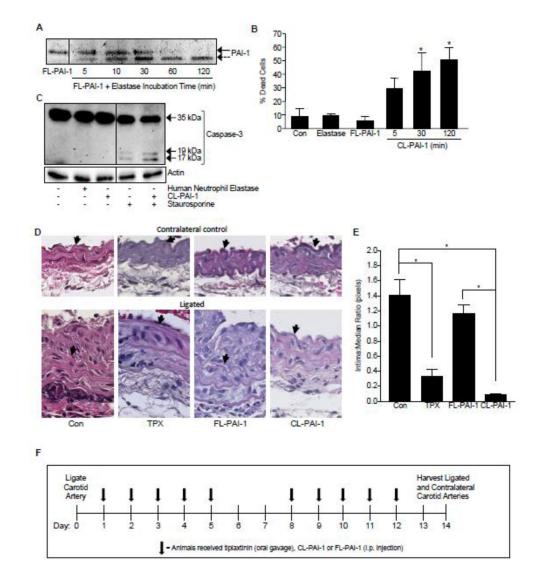


Figure 5. Elastase-cleaved PAI-1 promotes VSMC apoptosis while tiplaxtinin and CL-PAI-1 attenuate neointima formation

(A) Generation of CL-PAI-1 by incubation with human neutrophil elastase. Solid arrow indicates FL-PAI-1; dashed arrow denotes CL-PAI-1. (B) HuCASMCs treated with a final concentration of 10 nM FL-PAI-1 or CL-PAI-1 and a molar equivalent of elastase (as control) for 24 h were stained with ethidium homodimer and calcein AM to determine apoptotic index. Cells positive for ethidium homodimer were scored as apoptotic and normalized against calcein AM (total cells). Data are presented as mean \pm SEM; asterisks=p<0.05, n=3. (C) HuCASMCs were treated with CL-PAI-1 (40 nM) and/or staruosporine (5 μ M) as indicated for 24 h. Caspase-3 and actin (loading control) were assessed in whole cell lysates by western blot. (D) H&E stained cross-sections of contralateral control and ligated carotid arteries isolated from mice treated once daily with vehicle control (Con), 3 mg/kg tiplaxtinin (via oral gavage), 2 mg/kg FL-PAI-1 or CL-PAI-1 (both by intraperitoneal injection) for 14 days following ligation. Arrowheads indicate the internal elastic lamina. (E) Quantitation of the intima-to-media ration for the indicated

ligated carotid artery groups. Data plotted represent the mean \pm SEM; asterisks=p<0.05, n=4. (F) Time line of carotid ligation experiments; arrows indicate days of treatment with the indicated reagents.

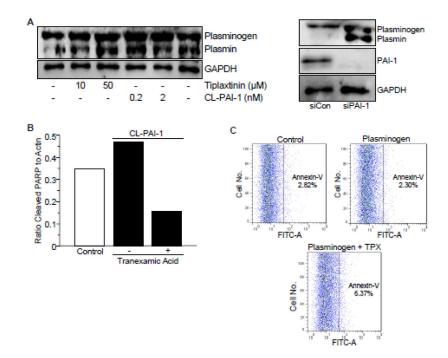


Figure 6. Cleaved PAI-1 promotes VSMC apoptosis in a plasmin-dependent manner (**A**) HuCASMCs were treated with tiplaxtinin or CL-PAI-1 for 6 h (left) or with the indicated siRNAs for 48 h (right). Plasminogen, plasmin, PAI-1 and GAPDH (loading control) were evaluated in cell lysates by western blotting. (**B**) HuCASMCs were incubated with 100 μ M tranexamic acid for 30 min prior to the addition of 40 nM CL-PAI-1. Cleaved PARP and actin (loading control) were assessed in cell lysates. (**C**) RASMCs were pretreated with 1.25 μ M plasminogen for 30 min prior to addition of tiplaxtinin (10 μ M, final concentration) for 24 h. Cells were collected, stained with Annexin-V and analysed by FACS. TPX=tiplaxtinin; CL-PAI-1=cleaved PAI-1

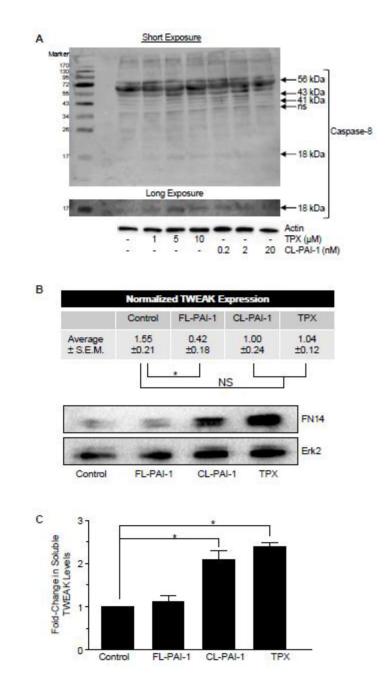


Figure 7. Upregulation of caspase-8 activation, FN14 and soluble TWEAK in response to cleaved PAI-1 and tiplaxtinin

(A) HuCASMCs were incubated with tiplaxtinin or CL-PAI-1 in the concentrations indicated for 24 h. Whole cell extracts were separated by gel electrophoresis and blots probed with antibodies to caspase-8 and actin (loading control). Bands: 56 kDa=full-length caspase-8; 43, 41 kDa= cleaved caspase-8; 18 kDa= cleaved cleaved caspase-8; ns=non-specific band. (B) HuCASMCs were treated with 40 nM FL-PAI-1, 40 nM CL-PAI-1 or 10 μ M tiplaxtinin for 24 hours, extracted and western blots developed using antibodies to TWEAK (depicted in table in **B** as mean ± SEM normalized to Erk2 expression, n=3) and FN14 using Erk2 as a loading control; NS=not significant. (C) Histogram is a plot of the

 $\label{eq:mean} \begin{array}{l} \text{mean} \pm \text{SEM} \text{ of the conditioned media levels of soluble TWEAK (relative to Ponceau S-stained total protein), n=4. TPX=tiplaxtinin; FL-PAI-1=full length PAI-1; CL-PAI-1=cleaved PAI-1 \end{array}$

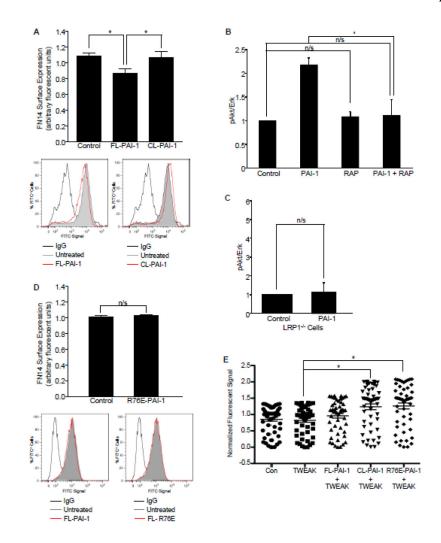


Figure 8. Full-length PAI-1 down-regulates FN14 in an LRP1-dependent manner and protects against TWEAK-stimulated apoptosis

(A) HuCASMCs were treated with 40 nM FL- PAI-1 or CL-PAI-1 for 24 h. Surface FN14 expression was assessed by FACS. Graphed in (A) is a plot of the mean \pm SEM; asterisks=p<0.05, n=4. Below the graph is a representative FACS histogram. n/s indicates no statistical significance. (B) PAI-1 induction of pAkt is LRP1-dependent. RASMCs were serum-starved for 48 hours before a 5 minute pretreatment with 5 µg/ml RAP prior to PAI-1 stimulation for 30 minutes. Cells were lysed and pAkt/Erk levels assessed by western blot analysis; pAkt levels were normalized to Erk1 and represented as a mean ± SEM, n=3. (C) LRP1^{-/-} cells were treated with vehicle alone or 40 nM PAI-1 for 30 minutes; lysates were analyzed by western blot analysis. pAkt levels were normalized to total Erk1 levels. Data is represented as the mean \pm SEM, n=3. Asterisks = p <0.05. n/s indicates no statistical significance. (D) RASMCs were incubated with 40 nM R76E-PAI-1 for 24 h, a recombinant PAI-1 mutant deficient in LRP1-binding. FACS was used to quantify surface FN14 expression. Graphed in (**D**) is a plot of the mean \pm SEM; asterisks=p<0.05, n=4. n/s indicates no statistical significance. Below the graph is a representative FACS histogram. (E) HuCASMC were treated with 40 nM FL- PAI-1, CL-PAI-1 or 40 nM R76E-PAI-1 and 250 ng/ml soluble TWEAK, or TWEAK alone, for 6 h. Cells were collected, lysed and an

aminomethylcoumarin (AMC)-tagged caspase-3 target (Z-DEVD) added. Fluorescence emission was monitored using a microplate reader every hour for 24 h. Data plotted represents the mean fluorescent intensity over a 24-h period ± SEM; asterisks=p<0.05 using a repeated-measures ANOVA. TPX=tiplaxtinin; FL-PAI-1=full length PAI-1; CL-PAI-1=cleaved PAI-1; R76E-PAI-1=LRP1-binding deficient mutant recombinant PAI-1 protein

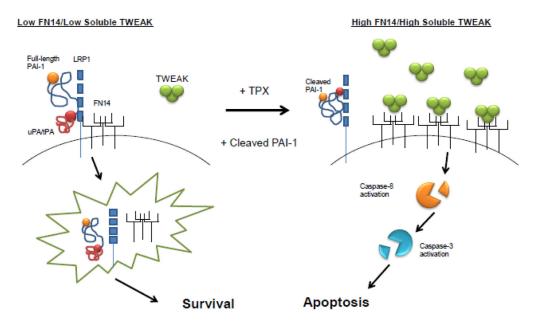


Figure 9. Hypothetical model

Current findings suggest that PAI-1 regulates VSMC survival and potentiates neointimal formation. Full-length PAI-1 binds to, and inhibits, uPA attenuating the conversion of plasminogen to plasmin while simultaneously down-regulating surface FN14 receptor expression through LRP1-mediated endocytosis, promoting a pro-survival, pro-stenotic phenotype. When the conformational pools of PAI-1 shift to increased levels of cleaved PAI-1 (either by tiplaxtinin or exogenous addition), conversion of plasminogen to plasmin is amplified. Membrane-bound and soluble TWEAK expression is increased stimulating, thereby, FN14-TWEAK signaling leading to VSMC apoptosis and reduced neointimal hyperplasia.