



The plasminogen activator inhibitor-1 paradox in cancer: a mechanistic understanding

Marta Helena Kubala^{1,2} · Yves Albert DeClerck^{1,2,3}

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Abstract

The paradoxical pro-tumorigenic function of plasminogen activator inhibitor 1 (PAI-1, aka Serpin E1) in cancer progression and metastasis has been the subject of an abundant scientific literature that has pointed to a pro-angiogenic role, a growth and migration stimulatory function, and an anti-apoptotic activity, all directed toward promoting tumor growth, cancer cell survival, and metastasis. With uPA, PAI-1 is among the most reliable biomarkers and prognosticators in many cancer types. More recently, a novel pro-tumorigenic function of PAI-1 in cancer-related inflammation has been demonstrated. These multifaceted activities of PAI-1 in cancer progression are explained by the complex structure of PAI-1 and its multiple functions that go beyond its anti-fibrinolytic and anti-plasminogen activation activities. However, despite the multiple evidences supporting a pro-tumorigenic role of PAI-1 in cancer, and the development of several inhibitors, targeting PAI-1, has remained elusive. In this article, the various mechanisms responsible for the pro-tumorigenic functions of PAI-1 are reviewed with emphasis on its more recently described contribution to cancer inflammation. The challenges of targeting PAI-1 in cancer therapy are then discussed.

Keywords Plasminogen activator inhibitor 1 · PAI-1 · Protease inhibitor · SerpinE1 · Tumor microenvironment · Inflammation

1 Introduction: the PAI-1 paradox

The seminal observation made by Riech's laboratory in 1973 that urokinase plasminogen activator (uPA) was the most robustly induced enzymatic activity in Rous Sarcoma Virus-transformed chicken cells [1] brought significant attention to the potential pro-tumorigenic role of plasminogen activation in cancer. Its importance was further underlined when multiple clinical studies revealed that cancer patients whose tumors have a high content of uPA and its receptor uPAR have worse clinical outcomes [2–4].

It was thus expected that because of its primary inhibitory activity on uPA, plasminogen activator inhibitor 1 (PAI-1), a member of the serine protease inhibitor (Serpin) family, would

have an anti-tumor effect, and it came as a surprise when these same studies and others reported high levels of PAI-1 in tumors that were associated with a poor rather than a favorable clinical outcome and that were less responsive to chemotherapy [5–11].

2 Complex and dynamic structure of PAI-1

Central to discovering the mechanisms supporting the pro-tumorigenic role of PAI-1 is the understanding of its complex and dynamic structure that explains its pleiotropic biological function (Fig. 1). PAI-1 is a 45 kDa single-chain glycoprotein of 379–381 residues with 9 α helices and 3 β sheets [12]. A unique structural aspect of PAI-1 and other members of the Serpin family is the presence of a 26 (Ser 343–Arg 368) amino acid reactive center loop (RCL) in the C-terminal region of the molecule that contains a proteolytic cleavage (P1-P1', Arg 346–Met 347) for uPA. Depending on the state of the RCL, PAI-1 exists in three forms. An active form that has the RCL in an helical configuration exposed at the surface of the molecule, a latent form unable to interact with uPA with the intact RCL as a strand internalized (analogy to the blade of a jack-knife) in the β -sheet A, and a cleaved form where the P1' Met residue of the RCL is exposed and the N-terminal portion of

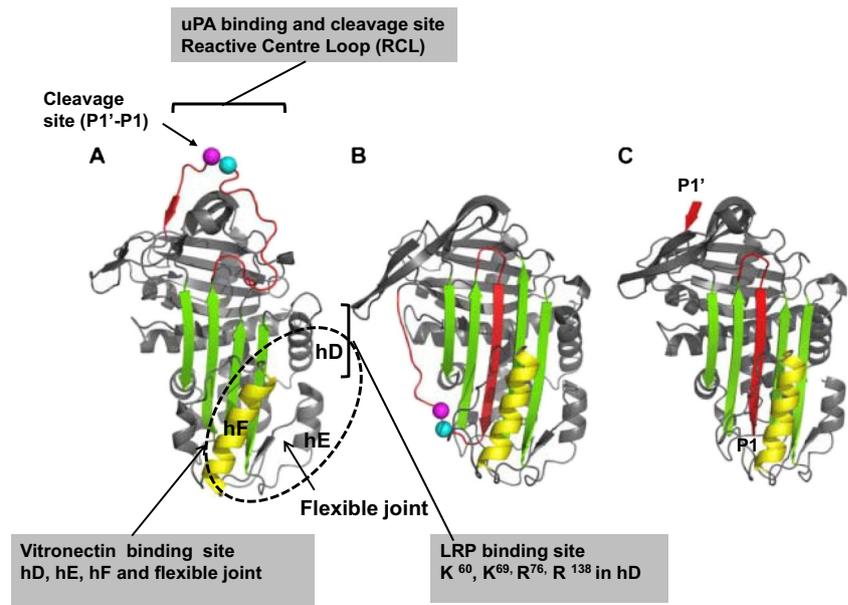
✉ Yves Albert DeClerck
declerck@usc.edu

¹ Division of Hematology, Oncology and Blood and Bone Marrow Transplantation, Department of Pediatrics, University of Southern California, Los Angeles, CA 90033, USA

² The Saban Research Institute of Children's Hospital, Los Angeles, CA 90027, USA

³ Department of Biochemistry and Molecular Medicine, University of Southern California, Los Angeles, CA 90033, USA

Fig. 1 Structure and function of PAI-1. The reactive center loop (RCL) binds to uPA and blocks uPA-generated plasmin. It is “internalized” as an additional strand (s4A in red) in the β -sheet A (in purple) upon latency and cleaved at P1-P1' by uPA. In blue, β sheet B whose position is modified upon latency. The helix D (hD) contains basic aa, which are involved in binding to LRP1. The region containing helices hD, hE, and hF (flexible joint) binds to the somatomedin B (SMB) domain in the N-terminal domain of vitronectin, preventing interaction with integrin. Adapted from [12]



the RCL containing the P1 residue remains inside the β sheet A.

Three domains binding to distinct proteins have been identified in PAI-1. The uPA binding domain is located in the RCL and binding of uPA to the RCL triggers its cleavage at the P1-P1' site, followed by the internalization of the N-terminal portion of the RCL containing the P1 site and uPA being translocated to the opposite pole of the molecule [13]. A second domain located in a flexible joint region between helix hD and hF spanning from residue 13 to 147, binds to the somatomedin B domain of vitronectin [14, 15]. Binding of PAI-1 to the somatomedin B domain of vitronectin stabilizes PAI-1, delaying its internalization in a latent form and increasing its anti-uPA ability. The binding of PAI-1 to vitronectin also masks the adjacent RGD binding site for αv integrins and inhibits cell adhesion to vitronectin [16, 17]. A third domain located in the N-terminal helix hD with a critical Lys 69 residue binds PAI-1 alone or complexed with uPA and tissue (t)PA to the endocytosis lipoprotein receptor proteins (LRP)-1 and (LRP)-2. Binding of PAI-1 to uPA at the cell surface *via* the attachment of uPA to its receptor (uPAR) triggers an interaction with LRP1 which leads to the intracellular internalization of the entire complex [18, 19]. A: Active PAI-1. B; Latent PAI-1. C; Cleaved PAI-1. Adapted from [20].

3 PAI-1 as a biomarker guiding cancer therapy

Numerous studies published in the late 1990s and early 2000s reported that PAI-1 is overexpressed by most human cancer cell lines—in particular, those derived from solid tumors—and in most human neoplasms. Its expression positively correlates with poor clinical outcome in patients with breast,

ovarian, bladder, colon, and non-small cell lung cancers [8, 21–28].

As a result of these observations, the prognostic value of the tissue levels of uPA and PAI-1 was tested in a large multinational (European Organisation for Research and Treatment of Cancer [EORTC], American Society of Clinical Oncology [ASCO], and Arbeitsgemeinschaft Gynäkologische Onkologie [AGO]) prospective clinical trial in 8377 patients with breast cancer without node involvement. This study, in which patients with high tumor tissue levels of uPA and/or PAI-1 determined by ELISA were randomized to 6 courses of chemotherapy or to observation, not only validated the long-term prognostic impact of uPA/PAI-1 levels but also demonstrated the benefit from adjuvant chemotherapy in the high-uPA/PAI-1 group [24].

4 PAI-1 polymorphism and cancer prediction

The PAI-1 (SERPINE1) gene is located on chromosome 7 (7q21.3-q22) and a polymorphism in the promoter of the gene has been described at position -675. This polymorphism known as 4G consists of deletion of a guanosine nucleotide. The 4G/4G homozygous polymorphism has a stronger binding for transcription factors whereas the 5G polymorphism allows the binding of repressor elements. As a result, higher levels of PAI-1 and a slight increase in deep venous thrombosis are observed in individuals with the 4G genotype [29]. These discoveries led to a series of clinical studies examining whether the 4G/5G polymorphism in patients with cancer could predict a more severe outcome in association with higher levels of PAI-1. Whereas several studies in breast, hepatocellular carcinoma, testicular, endometrial, and oral

cancer indicated a correlation between 4G/5G polymorphism, higher levels of PAI-1, and overall poor clinical outcome [30–35], other studies in breast, renal cell, prostate, and colorectal cancers have failed to show such correlation [31, 36–40]. A meta-analysis of 10 publications on 4G/5G polymorphism in breast cancer revealed a 39% increase in the incidence of breast cancer in populations with the 4G/4G homozygous phenotype. The study also revealed an association between the 5G/5G genotype and the presence of lymph node metastasis. No differences between genotype and histological grade and overall survival were identified [41]. The issue of the role of PAI-1 gene polymorphism in cancer is thus not entirely clear.

5 PAI-1 and the hallmarks of cancer

Using the text mining Cancer Hallmark Analytical Tool (<http://chat.lionproject.net>) [42], 3739 articles and abstracts with PAI-1 were assigned to one of the 10 Hallmarks of Cancer [43] with a pre-eminent role in five hallmarks, sustaining proliferative signals, resisting cell death, angiogenesis, invasion and metastasis, and tumor-promoting inflammation (Fig. 2). We discuss here the role of PAI-1 in these five hallmarks with an emphasis on the PAI-1 interactive domain involved and the mechanisms identified (Table 1).

6 Role of PAI-1 in sustaining proliferative signals

Some studies have shown that PAI-1 has a growth stimulatory activity leading to G1-phase to S phase cell-cycle progression through cyclin D3/cdk4/6 upregulation in cancer cells [44]. However, other studies in human breast cancer cell lines have indicated an inhibitory function of PAI-1 on anchorage-dependent and independent

proliferation [77]. Whether PAI-1 directly affects tumor cell cycle remains controversial as long as no precise molecular mechanism is identified.

PAI-1 also has an indirect modulatory effect on cancer cell growth *via* its anti-fibrinolytic activity and modulatory activity on cell adhesion and uPA/uPAR activity. By inhibiting fibrinolysis, PAI-1 maintains the activity of thrombin which interacts with protease-activated receptors (PAR) in tumor cells and enhances proliferation *via* PAR activation [48]. Thrombin interacting with PAR also transcriptionally upregulates PAI-1 [49]. By promoting detachment from vitronectin and adhesion to fibronectin, PAI-1 stimulates fibronectin-dependent cell growth [45] and by downregulating uPA/uPAR interaction with EGFR, PAI-1 inhibits EGF-dependent cell growth [50]. Studies have also shown that PAI-1 plays a role in senescence [46]. PAI-1, a target of p53 and TGF- β , is upregulated in aging fibroblasts and inhibits the release and bio-availability of many growth factors. Suppression of PAI-1 in aging fibroblasts stimulates proliferation and allows them to bypass senescence. PAI-1 may also stimulate dormancy through uPAR and uPA inhibition. In breast, prostate, melanoma, and fibrosarcoma cell lines, the ERK/p38 activity ratio determines whether *in vivo* cells will proliferate or go into dormancy. A high ERK/p38 ratio favors tumor growth, whereas a low ERK/p38 ratio induces dormancy. The ERK/p38 ratio is affected by pericellular uPA activity. A high-uPA and uPAR activity stimulates $\alpha 5\beta 1$ -integrin and activates ERK1/2 resulting in a high ERK/p38 ratio. By contrast, the inhibition of uPA/uPAR activity by PAI-1 reverses the ratio inhibiting growth and promoting cell dormancy [47]. PAI-1 is therefore more a modulator of cancer cell proliferation, having complex activities involving multiple interactions with uPA, uPAR, integrins, extracellular matrix (ECM) proteins that have indirect effects on the signaling activity of thrombin, growth factor receptors, and integrins.

Fig. 2 Role of PAI-1 in cancer progression. The doughnut diagram represents the proportion of publications assigned to each indicated hallmark and was obtained using the Cancer Hallmarks Analytics Tool [42] (<http://chat.lionproject.net>). The numbers inside the doughnut represent the number of publications

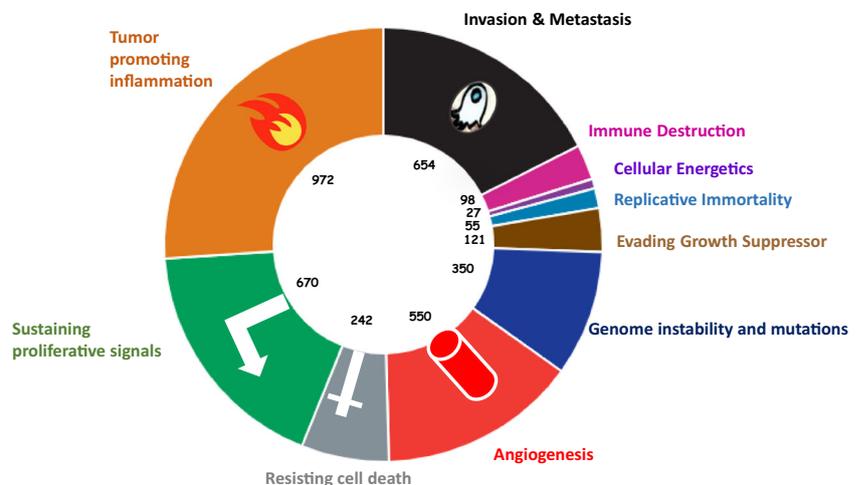


Table 1 Multifaceted activities of PAI-1 in human cancer

Hallmark	Effect	PAI-1 domain	Mechanism
Sustaining proliferative signals	Increases transition from G0/G1 to S phase	Unknown	Upregulation of cyclin E and cyclin E complexes with CDK with decrease in p53, p21, and p27 [44]
	Modulates cell adhesion	Vitronectin binding	Increases adhesion/proliferation to fibronectin [45]
	Promotes senescence and dormancy	uPA binding (RCL)	Altering ERK/p38 ratio <i>via</i> inhibition of uPA/uPAR [46, 47]
	Inhibition of fibrinolysis	uPA binding (RCL)	Increase in PAR stimulation by thrombin; PAR activation upregulates PAI-1 [48, 49]
	Inhibits EGFR signaling	uPA/uPAR/PAI-1 complex	Inhibits uPA/uPAR interaction with EGFR [50]
Resisting cell death	Decreases intrinsic apoptosis	uPA binding (RCL)	Intracellular inhibition of caspase 3 [51]
	Induces anoikis	Vitronectin binding domain	Inhibition of cell adhesion to vitronectin, increases cell migration and adhesion to fibronectin (double edge) [52]
	Decreases apoptosis	uPA binding (RCL)	Decrease in FasL cleavage and shedding by plasmin, inhibition of FasL-mediated apoptosis in tumor cells [53, 54]
Angiogenesis	Protects from apoptosis	LRP-1 binding	Stimulates c-Jun/ERK and increases expression of Bcl-2 and Bcl-XL [52]
	Increases EC survival	uPA binding (RCL)	Decrease in FasL cleavage and shedding by plasmin, inhibition of FasL-mediated apoptosis in EC [55]
	Inhibits EC adhesion to vitronectin	Vitronectin binding domain	Stimulation of EC migration from vitronectin toward fibronectin [56]
	Increases EC organization in fibrin matrix	uPA binding (RCL)	Increase in EC adhesion to fibrin and IL-8 production [57, 58]
Invasion and metastasis	Modulates tumor cell migration	uPA binding and vitronectin binding	Promotes cell detachment from vitronectin and prevents excessive degradation of ECM proteins [59]
	Increases <i>in vivo</i> metastasis	Unknown	Not clearly defined [60–67]
	Decreases <i>in vivo</i> metastasis	Unknown	Not clearly defined [68–70]
	Inhibition of fibrinolysis	uPA binding (RCL)	Promotes NET [71, 72]
Tumor promoting inflammation	Increases monocyte/macrophage migration	LRP binding	Recruits monocytes to tumor site [73–75]
	Promotes macrophage M2 polarization	uPA binding (RCL)	Activates p38/NFκB/IL-6/STAT3 [76]

7 Role of PAI-1 in resisting tumor cell death

The anti-apoptotic activity of PAI-1 has been extensively described and the mechanisms have been well characterized [78]. By inhibiting cell adhesion to vitronectin, PAI-1 can have a pro- and anti-apoptotic effect. By promoting cell detachment, it stimulates anoikis, but as cells detach from vitronectin and migrate toward other ECM proteins, they can resist apoptosis [52]. As a serpin, PAI-1 inhibits intracellular caspase 3, protecting tumor cells from chemotherapy-induced apoptosis [51]. Extracellular PAI-1 also inhibits the cleavage and shedding of FasL by plasmin at the surface of tumor cells protecting them from FasL-mediated and chemotherapy-induced cell death [53, 54]. By interacting with LRP-1, PAI-1 induces c-Jun/ERK signaling driving an increase in the production of anti-apoptotic proteins like Bcl 2 and Bcl-X_L [52].

8 Role of PAI-1 in angiogenesis

PAI-1 is a stimulator of angiogenesis by promoting migration, survival, and proliferation in endothelial cells (EC). This pro-angiogenic activity stems from its protease-inhibiting and vitronectin-binding functions [55, 79]. As an inhibitor of plasminogen activation, PAI-1 protects EC from FasL-dependent extrinsic apoptosis [55]. Through its ability to bind to the somatomedin B domain of vitronectin, PAI-1 interferes with the binding of EC to vitronectin promoting EC detachment from vitronectin and migration toward a fibronectin-rich but less vascularized tumor stroma [56]. Through its anti-fibrinolytic activity, PAI-1 increases fibrin deposition which promotes EC organization [57] and the release of angiogenic proteins like interleukin (IL)-8 [58].

9 Role of PAI-1 in invasion and metastasis

Many *in vitro* experiments have demonstrated that PAI-1 promotes tumor cell migration. By preventing the adhesion of tumor cells to vitronectin, PAI-1 stimulates their migration toward other ECM substrates like fibronectin [80]. By inhibiting uPAR-bound uPA, PAI-1 also prevents excessive pericellular degradation of ECM proteins necessary for cell adhesion and migration [59]. PAI-1 thus plays a role of the modulator of cell migration by both promoting cell detachment from some ECM proteins like vitronectin and by preventing excessive pericellular ECM degradation. By contrast, *in vivo* studies in metastasis have reported conflicting results, with some experiments suggesting a pro-metastatic effect [60–67] and others either an inhibitory effect [68–70] or an absence of effect [81] on metastasis [82]. The discordance in these data is still not well understood and illustrates the complexity of studies related to metastasis in mice [20]. A major problem in the interpretation of these studies is the fact that they are based on a variety of routes of tumor cell administration (experimental, subcutaneous, and orthotopic implantation), different mouse background (immunocompetent, immunodeficient, or transgenic mice), and different experimental approaches (pharmacological and genetic knock-down, genetic knock-out, overexpression). Whereas some studies use the macroscopic evaluation of metastasis, others have relied on a more precise microscopic detection of micrometastases. The amount of PAI-1 produced and its source (tumor cells vs. host cells) also play a critical role, with host-derived PAI-1 having a more critical role [83]. The discordance among the data reported also needs to take into consideration the differences between human and mouse PAI-1 and their plasminogen activation system [84]. Thus, although clinical and pre-clinical studies suggest that PAI-1 stimulates the formation of metastasis, in the absence of more mechanistic experiments, the precise contribution of PAI-1 to metastasis remains elusive.

One potential novel mechanism by which PAI-1 could stimulate the extravasation of circulating tumor cells (CTC) at the site of metastasis is *via* neutrophil extracellular traps (NET). These recently described structures made of DNA and proteolytic enzymes formed during infection could support the formation of metastasis [71]. By activating the formation of microthrombi through its anti-fibrinolytic activity, PAI-1 may stimulate the extravasation of CTC to metastatic sites in NET [72].

10 Role of PAI-1 in cancer inflammation

PAI-1 levels are elevated in non-malignant inflammatory conditions such as sepsis, metabolic syndromes (MetS), and arthritis [85–88]. Experimental *in vivo* studies performed in

animal models as well as in humans have shown that inflammatory cytokines like tumor necrosis factor- α (TNF- α) and IL-6 stimulate PAI-1 expression [89, 90], corroborating previously reported correlation between PAI-1 expression and inflammation [91]. These observations indicated that PAI-1 is a downstream consequence of inflammation and a contributor to the thrombotic and anti-fibrinolytic syndromes associated with inflammation and cancer as initially observed by Armand Trousseau in 1865 [92]. Adipocytokines like insulin, insulin-like growth factor-1, TNF- α , IL-6 and leptin/adiponectin, and other by-products of MetS such as glucose and cholesterol alter PAI-1 expression, not only in stromal cells but also in cancer cells to potentially favor invasion and metastasis [77]. However, more recent studies suggest that PAI-1 is not only a consequence of but also a contributor to inflammation. PAI-1 increases the production of IL-8 and leukotriene B4 in alveolar epithelial cells and promotes the migration of inflammatory cells [93]. PAI-1 stimulates the recruitment of fibrosis-inducing cells and macrophages [73–75] and the survival of T lymphocytes [94]. The contributory role of PAI-1 in cancer-related inflammation has only recently been explored. Our laboratory has recently demonstrated that through its interaction with LRP-1, tumor-derived PAI-1 promotes the migration of monocytes [76]. PAI-1 also influences the polarization of monocytes/macrophages toward an M2 pro-tumorigenic function through the activation of p38MAPK and NF κ B in macrophages and the transcriptional upregulation of IL-6. *Via* an autocrine loop, IL-6 activates signal transduction and activator of transcription (STAT) in monocytes leading toward an increase in the expression of arginase, IL-10, and CD163. Mutational analysis demonstrated that this latter function requires the C-terminal uPA-interacting domain and is independent of the N-terminal LRP-1 binding domain required for the stimulation of monocyte migration [76].

11 Targeting PAI-1

Considering the abundant scientific literature supporting a pro-tumorigenic role of PAI-1 in human cancer, it is surprising that there is still no definitive evidence that inhibition of PAI-1 could have any therapeutic effect in cancer patients. Numerous small molecule and antibody inhibitors of PAI-1 have been developed over the last 10 years [95] and several have been tested in preclinical animal models. Whereas they have shown efficacy in models of acute thrombosis, promoting rapid thrombus re-permeabilization, their activity in models of cancer progression and metastasis is less clear, with the demonstration of an inhibitory effect on tumor growth and metastasis in some cases and an absence of significant effect in other cases [20].

The first group of small inhibitors, SK-116 and SK-216, were shown to cause an almost 2-fold reduction in the number

Table 2 Roles of PAI-1 in cancer

Pro-tumorigenic activity	Established
Biomarkers of poor outcome	Established
Biomarker guiding therapy	Established
Prognostic value of G4/G5 polymorphism	Controversial
Role in carcinogenesis	Controversial
Role in cancer cell proliferation	Not fully understood
Role in dormancy	Suggested
Protective role in apoptosis	Established
Role in angiogenesis	Established
Role in cell migration	Established
Role in metastasis	Controversial
Role in cancer inflammation	Emerging

of small intestinal polyps in Min mice [96] and a 2-fold reduction in the growth of subcutaneous primary B16 melanoma tumors and metastasis in mice [67]. They were also shown to have preclinical activity in malignant pleural mesothelioma [97]. PAI-039, Tiplaxtinin, an indol-oxoacetic acid-based inhibitor extensively tested in acute thrombosis, has been shown to have anti-tumor activity in T47 bladder cancer and HeLa cell tumors in mice [98]. TM 5275 and 554, N-acylanthranilic acid derivatives were found active in some preclinical cancer models but not in others [99, 100]. However, none of these inhibitors have been tested in therapeutic clinical trials in cancer. The effectiveness of PAI-1 inhibitors has been limited by their lack of activity against the stable, vitronectin-bound form of PAI-1, their insufficient pharmacokinetic properties with a half-life of less than 2 h, and high therapeutic concentrations in the μM range [101] and their testing as a single agent. The need for a high therapeutic index is another important consideration since PAI-1 inhibition will require a chronic administration to prevent cancer metastasis or induce dormancy. The use of these inhibitors should be re-evaluated within the context of combination therapies, including immunotherapy, considering their recently identified role on macrophage recruitment and polarization in the TME.

12 Conclusion

More than 30 years have passed since the first reports of the expression of PAI-1 in cancer cells by Kretzmer and Vassali's laboratories [102, 103]. There is now an extensive literature that supports the pro-tumorigenic activity of PAI-1. The role and use of PAI-1 as a biomarker of cancer prognosis and response to therapy are well established. The mechanisms explaining most of the pro-tumorigenic activities of PAI-1 are well understood (Table 2). Targeting PAI-1 in cancer remains challenging, in part because of difficulties in the design of more effective inhibitors against active PAI-1 suitable for

chronic administration. Considering the important role of PAI-1 in the TME and in cancer-related inflammation, PAI-1 could have a function in immune escape and targeting PAI-1 in combination with immunotherapy should be an important aspect to investigate.

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