

# TIMP-3 as a therapeutic target for cancer

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*Ther Adv Med Oncol*

2019, Vol. 11: 1–17

DOI: 10.1177/  
1758835919864247

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**Abstract:** Tissue inhibitor of metalloproteinase-3 (TIMP-3), a secreted glycoprotein, plays an important role in carcinogenesis. It can bind to many proteinases to suppress their activity and thus protect the extracellular matrix from degradation. TIMP-3 may have many anticancer properties, including apoptosis induction and antiproliferative, antiangiogenic, and antimetastatic activities. This review summarizes the structure, proteinase inhibition ability, genetic and epigenetic regulation, cancer therapy potential, and contribution to cancer development of TIMP-3. Furthermore, in this review we discuss its potential as a biomarker for predicting cancer progression and the current state of drugs that target TIMP-3, either alone or in combination with clinical treatment. In conclusion, TIMP-3 can be a biomarker of cancer and a potential target for cancer therapy. This review article can serve as a basis to understand how to modulate TIMP-3 levels as a drug target of cancers.

**Keywords:** cancer therapy, extracellular matrix, matrix metalloproteinase, metastasis, tissue inhibitors of metalloproteinases-3

Received: 14 November 2018; revised manuscript accepted: 3 June 2019.

## Introduction

Despite recent improvements, cancer treatment remains associated with several challenges. Cancer development is a multifactorial and multistep process and involves several genetic and epigenetic regulations. In addition, cancer cells possess several unique characteristics, also known as the cancer hallmark, that confer the cells with resistance against the human immune system and cancer treatment. The cancer hallmark includes tumor-promoting inflammation, enabling replicative immortality, avoiding immune destruction, evading growth suppressors, sustaining proliferative signaling, deregulating cellular energetics, resisting cell death, genome instability and mutation, inducing angiogenesis, and activating invasion and metastasis.<sup>1,2</sup> Metastasis, which is the major cause of death among cancer patients, involves multiple processes including extracellular matrix (ECM) remodeling and degradation. Degradation of the ECM is required for tumor cell metastasis; this is achieved by several proteinases such as the plasmin system and particularly, the matrix metalloproteinases (MMPs).<sup>3–6</sup> MMPs are known to play an important role in the tissue invasion and metastasis of cancer cells. The tissue inhibitors of MMPs (TIMPs) are endogenous

inhibitors of MMPs, and regulation of MMPs by TIMPs is particularly important for the maintenance of the ECM. Disruption of the balance between the activities of MMPs and TIMPs during carcinogenesis may affect invasion and metastasis<sup>7–9</sup> and may worsen patient outcomes.<sup>10</sup> TIMP-3, a member of the TIMP family, is a 24-kDa secreted glycoprotein, and its gene is located on chromosome 22q12.1–q13.2. Knockout of the *TIMP-3* gene in mice resulted in increased MMP, a disintegrin and MMPs with thrombospondin motifs (ADAMTS) activity, and cartilage degradation, suggesting that reduced TIMP-3 levels may cause osteoarthritis.<sup>11</sup> In addition, the absence of TIMP-3 leads to poor cardiac remodeling and has been associated with myocardial infarction or hypertension.<sup>12,13</sup> In cancer studies, TIMP-3 plays an important role in the cancer hallmark by controlling cell death, angiogenesis, tumor inflammation, and tumor cell invasion and dissemination.<sup>14</sup> For instance, TIMP-3 restoration in cancer cells inhibits cell growth and promotes cell apoptosis.<sup>15,16</sup> In addition, TIMP-3 overexpression improves the sensitivity of osteosarcoma to clinical drug treatment through interleukin (IL)-6 inhibition.<sup>17</sup> TIMP-3 also acts as a potential antiangiogenesis agent by

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inhibiting endothelial cell tube formation.<sup>18</sup> Moreover, TIMP-3 can inhibit cancer cell migration, invasion, and metastasis *in vitro* and *in vivo*.<sup>19,20</sup> Clinical studies have reported reduced TIMP-3 expression in cases of several cancer types compared with normal controls;<sup>19–22</sup> the loss of TIMP-3 may lead to poor outcomes, including large tumor size, high tumor stage, and metastasis.<sup>23–25</sup> Herein, we review the structure and function of TIMP-3 and discuss its contribution to carcinogenesis and its potential in cancer therapy.

## TIMP-3

### *TIMP classification*

The TIMP family contains four members: TIMP-1, TIMP-2, TIMP-3, and TIMP-4. The molecular weight of the TIMPs is approximately 21 kDa, and they contain an N-terminal domain and a C-terminal domain. In contrast to that in TIMP-1 and TIMP-3, the C-terminal domain in TIMP-2 and TIMP-4 is negatively charged. Compared with the nonglycosylation of TIMP-2 and TIMP-4, TIMP-1 contains two N-glycosylation sites at Asn<sup>26</sup> and Asn,<sup>27</sup> and TIMP-3 contains a single N-glycosylation site in the C-terminal domain.<sup>28</sup> Glycosylated forms of TIMP-3 have higher affinity for glycan-bound MMPs and are protected from endocytosis and degradation.<sup>29</sup> Unlike other TIMPs, TIMP-3 is the only TIMP that binds firmly to the ECM after its secretion.<sup>30</sup> This binding is *via* the interaction of the N-terminal domain with heparan sulfate and sulfated glycosaminoglycans.<sup>31</sup>

### *Transcriptional regulation of TIMP-3*

The expression of TIMP-3 can be regulated by transcriptional regulation. Transcriptional regulation contains two major parts: the first part involves transcription factors and the transcription apparatus and the second part involves chromatin and its regulators.<sup>26</sup> Gene expression regulated by transcription factors is one of the most common transcriptional regulations. Transcription factors including Elf3, sp1, smad2, and smad4 have been reported to target on the promoter of TIMP-3 and regulated TIMP-3 expression.<sup>32–36</sup> Jobling *et al.* discovered that ETS transcription factor Elf-3 was expressed in human retinal pigment epithelium (RPE) cell lines. Transfection of Elf3a and Elf3b overexpression vector increased promoter activity of TIMP-3.<sup>32</sup>

TIMP-3 promoter contains four sp1 binding sites in the region near the transcription start site.<sup>35</sup> Zerrouqi *et al.* indicated that P14ARF increased expression of TIMP-3 in human glioblastoma cell line is sp1 dependent. Knockdown of sp1 by siRNA suppressed TIMP-3 promoter activity that is enhanced by P14ARF.<sup>34</sup> Other studies also demonstrated that sp1 regulated TIMP-3 promoter transcription activity *via* the ERK pathway.<sup>33,35</sup> Treatment of ERK inhibitor decreased binding ability of sp1 to DNA.<sup>35</sup> TIMP-3 is also a target for Smad pathway mediated by transforming growth factor (TGF)- $\beta$ . Qureshi *et al.* suggested that the transcription factors Smad2 and Smad4 must bind to the promoter of TIMP-3 in the presence of TGF- $\beta$ .<sup>36</sup> In addition, TIMP-3 expression can also be regulated by histone modification such as histone acetylation and histone methylation. Shinojima *et al.* used chromatin immunoprecipitation and showed that transcriptional repression of TIMP-3 was associated with increased H3K27me3 and decreased H3K9ac histone marks at TIMP-3 promoter.<sup>37</sup> Many proteins have also been reported to be involved in the process of histone modification. HDAC9 is one of the histone deacetylases (HDACs) that has been indicated to suppress TIMP-3 *via* promoter histone hypoacetylation.<sup>38</sup> KDM1A, also known as LSD1, caused TIMP-3 repression through H3K4me2 demethylation at TIMP-3 promoter.<sup>39</sup> The enhancer of zeste homolog 2 (EZH2), which has histone methyltransferase activity, is known to reduced TIMP-3 expression by catalyzing H3K27me3.<sup>40</sup>

### *MMP inhibitory activity of TIMP-3*

TIMPs are endogenous inhibitors of MMPs and exhibit marked antiproteinase activity against MMPs, ADAMs, and ADAMTSs.<sup>41</sup> TIMPs can use the N-terminal region to bind to the catalytic domain of MMPs to inhibit their activity and form a stable bond with the C-terminal hemopexin domain of proMMPs *via* the C-terminal region.<sup>42</sup> However, the extent of MMP inhibition differs between each TIMP; TIMP-1 strongly inhibits MMP-9 but poorly inhibits MT1-MMP, MT3-MMP, MT5-MMP, and MMP-19,<sup>30</sup> and TIMP-2 strongly inhibits MMP-2 and can inhibit other MMP members. TIMP-1, TIMP-2, and TIMP-4 inhibit only a few ADAMs.<sup>43–45</sup> In addition, TIMP-2 can form a ternary complex composed of TIMP-2-pro-MMP-2-MT1-MMP, which resulted in the activation of pro-MMP-2.<sup>30</sup> TIMP-4 can also form a TIMP-4-pro-MMP-2-MT1-MMP

complex, but unlike TIMP-2, leading to inhibit the activation of pro-MMP-2 *via* inhibition of MT1-MMP.<sup>46</sup> TIMP-3 can form a similar terminal complex to inhibit pro-MMP-2 activation. Knockout of TIMP-3 in cell promoted activation of pro-MMP-2 mediated by MT1-MMP.<sup>47</sup> In contrast to other members of the TIMP family with limited inhibitory activity for ADAMs, TIMP-3 can effectively inhibit ADAM10, ADAM12, ADAM17, ADAM28, ADAM33, ADAMTS-1, ADAMTS-2, ADAMTS-4, and ADAMTS-5.<sup>30</sup> For instance, the ECM protein-degrading activity of ADAM12 can only be blocked by TIMP-3, but not by TIMP-1, TIMP-2, and TIMP-4.<sup>48</sup> In addition, TIMP-2 and TIMP-3 inhibit the aggrecanase activity of ADAMTS1, but TIMP1 and TIMP4 have no significant inhibitory effect.<sup>49</sup>

#### *TIMP-3 in cancer*

Although TIMPs may have substantial roles in cancer, the mechanisms and outcomes for each TIMP vary. For example, high TIMP-1 expression has been reported to be associated with poor prognosis in most cancers. In an *in vitro* study, TIMP-1 expression was found to promote cancer cell survival in acute myeloid leukemia (AML) and melanoma.<sup>50,51</sup> Clinical studies have shown that a high level of TIMP-1 is associated with a shortened relapse-free and cancer-specific survival in endometrial carcinoma<sup>52</sup> and poor overall survival in laryngeal squamous cell carcinoma.<sup>53</sup> Both increased and decreased expressions of TIMP-2 and TIMP-4 have been reported to be associated with greater cancer risk. Overexpression of TIMP-2 has been reported to increase proliferation of choriocarcinoma cells.<sup>54</sup> By contrast, TIMP-2 inhibits angiogenesis *via* an MMP-independent mechanism.<sup>55</sup> TIMP-4 is expressed in some cancers; however, its role in carcinogenesis remains unclear. In a recent study, TIMP-4 was found to regulate stemness in cervical cancer by enriching tumor progenitor cells.<sup>56</sup> Studies in human cancer tissues have suggested that TIMP-3 may play a tumor suppressive role, and that TIMP-3 gene expression is downregulated in brain tumors, esophageal adenocarcinoma, gastric adenocarcinoma, clear cell renal cell carcinoma, meningiomas, and pancreatic endocrine tumors.<sup>57–61</sup> In an *in vitro* study, TIMP-3 expression was observed to induce cancer cell apoptosis in cervical cancer, fibrosarcoma, and breast cancer cell lines.<sup>62</sup> In an animal model, TIMP-3-deficient prostate tumors

showed increased expression of inflammation markers, including monocyte chemoattractant protein (MCP) 1, COX2, tumor necrosis factor (TNF)- $\alpha$ , and IL-1 $\beta$ .<sup>63</sup> In addition, TIMP-3 has been widely described as a potential angiogenesis inhibitor. A study reported that purified and renatured TIMP-3 inhibits angiogenesis, as revealed by the CAM assay.<sup>64</sup> Moreover, TIMP-3 can inhibit the migration and invasiveness of cancer cells *in vitro*.<sup>65–67</sup> By contrast, some reports have suggested that TIMP-3 promotes cancer. Higher *TIMP-3* mRNA levels were observed in the stroma of head and neck cancer cells than in the normal epithelial cells, and high levels of TIMP-3 have been shown to be associated with a significant reduction in the overall survival rate.<sup>68</sup> Our previous study also revealed that plasma TIMP-3 is a potential biomarker of the tumor stage in patients with oral squamous cell carcinoma.<sup>21</sup> *TIMP-3*-knockout mice demonstrated resistance to developing breast and liver cancer despite the increase in their inflammatory response.<sup>69,70</sup>

#### **Genetic and epigenetic regulation of TIMP-3 in cancer**

##### *Gene polymorphism*

Gene expression may be regulated by gene polymorphisms, which are variations in the DNA sequence resulting from a >1% nucleotide change within a population.<sup>71,72</sup> Polymorphisms of *TIMP-3* have been reported in many cancers including adenocarcinoma, bladder cancer, breast cancer, hepatocellular carcinoma, oral cancer, and prostate cancer.<sup>23,73–77</sup> The *TIMP-3* polymorphic rs9862 allele is associated with increased plasma levels of TIMP-3 and higher risk of oral cancer than with the wild-type allele.<sup>23</sup> In adenocarcinoma of the gastroesophageal junction, polymorphisms (rs130274, rs715572, rs1962223, and rs5754312) in *TIMP-3* and polymorphism rs9862 in the *TIMP-3* promoter are associated with survival.<sup>77</sup> Moreover, rs8136803 (TT) is associated with decreased disease-free survival in breast cancer patients.<sup>78</sup>

##### *DNA methylation and histone modification*

Loss or downregulation of TIMP-3 expression has been linked to promoter hypermethylation of the *TIMP-3* gene in several types of cancer including esophageal adenocarcinoma, head and neck squamous cell carcinoma, ovarian cancer, and

pancreatic endocrine tumors.<sup>27,61,79</sup> Hypermethylation in the promoter region usually causes transcriptional silencing because it affects the ability of the transcription factor to bind to the target gene. *TIMP-3* methylation in the sp1 binding site and the TATA box of the promoter are associated with low expression of *TIMP-3* protein in gastric cancer cell lines.<sup>80</sup> DNA methyltransferases (DNMTs) are key enzymes causing gene methylation, and dysregulation of DNMTs has been reported in tumorigenesis.<sup>81</sup> Knockout of both DNMT1 and DNMT3B in colorectal carcinoma cell lines caused gene demethylation of *TIMP-3*, which resulted in recovery of *TIMP-3* mRNA expression.<sup>82</sup> *TIMP-3* expression is also regulated by ten–eleven translocation 1 (TET1), a dioxygenase involved in cytosine demethylation. TET1 can maintain the expression of *TIMP-3* by inhibiting its methylation and, thus, suppressing cancer cell invasion.<sup>83</sup> Loss of *TIMP-3* expression can also be regulated by histone H3K27 methylation *via* upregulation of the EZH2 in non-small cell lung cancer (NSCLC).<sup>40</sup> In addition, *TIMP-3* expression was suppressed by KDM1A, a histone demethylase that removes H3K4me2 from *TIMP-3* promoter and promotes tumor cells invasion in NSCLC.<sup>39</sup> In prostate cancer, treatment with histone methylation inhibitor 3-deazaneplanocin A and trichostatin A restored expression of *TIMP-3*.<sup>37</sup>

#### MicroRNA and long noncoding RNAs

Accumulating evidence suggests that microRNAs (miRNAs) and long noncoding RNAs (lncRNAs) affect cancer development.<sup>84</sup> MiRNAs involved in the post-transcriptional regulation of *TIMP-3*. The mechanism of MicroRNA to suppress gene expression is by controlling mRNA stability and translation through base pairing to the 3' untranslated region (3'-UTR). *TIMP-3* regulation by miRNAs including miR-17-3p, miR-17-5p, miR-21, miR-21-5p, miR-181b, miR-191, miR-221, miR-222, and miR-373 has been widely reported in different cancers (Table 1).<sup>85–101</sup> For instance, mature miR-17-5p and the passenger strand miR-17-3p promoted prostate cancer growth and invasion by targeting *TIMP-3*.<sup>85</sup> Upregulation of miR-21 was correlated with decreased *TIMP-3* expression in patients with breast cancer, cervical cancer, cholangiocarcinoma, and pancreatic ductal adenocarcinoma.<sup>87,88,90,91</sup> In an *in vitro* analysis, miR-373 could induce esophageal squamous cell carcinoma cell migration and invasion by inhibiting *TIMP-3* expression.<sup>101</sup> In an *in vivo*

study, miR-181b upregulation by TGF- $\beta$  reduced *TIMP-3* expression, and transfection with anti-miR-181b in the hepatocellular carcinoma cell line SK Hep-1 suppressed tumor growth in nude mice.<sup>94</sup> Some studies have suggested that miRNAs targeting *TIMP-3* are also crucial in cancer resistance development. For instance, miR-21-5p was upregulated in patients with gastric cancer and induced drug resistance to doxorubicin by targeting *TIMP-3*.<sup>92</sup> Upregulation of miR-221 in oral squamous cell carcinoma cells increases resistance to doxorubicin by silencing *TIMP-3* expression.<sup>97</sup> Moreover, miR-221 inhibition enhances the sensitivity of human oral squamous cell carcinoma cells to doxorubicin by upregulating *TIMP-3* expression<sup>98</sup> (Figure 1). lncRNAs are RNA transcripts that are longer than 200 nucleotides. Accumulating evidence indicates that lncRNAs play critical roles in tumorigenesis through various mechanisms such as transcriptional, post-transcriptional, and epigenetic regulation.<sup>102–105</sup> The lncRNA BC032913 enhances *TIMP-3* expression and inhibits nuclear  $\beta$ -catenin expression, thus suppressing the migration, invasion, and metastatic potential of colorectal cancer cells.<sup>106</sup> The lncRNA DANCR suppresses *TIMP-3* expression by increasing the binding ability of EZH2 and H3K27me3 to the *TIMP-3* promoter in prostate cancer.<sup>107</sup>

#### Function of *TIMP-3* in cancer

*TIMP-3* functions as a tumor suppressor gene in many types of cancer. It exerts anticancer effects through mechanisms such as the inhibition of cell proliferation, the induction of apoptosis, the inhibition of drug resistance of cancer and inhibition of angiogenesis, migration and invasion, cancer metastasis, and epithelial–mesenchymal transition (Figure 2). In this section, we focus on the aforementioned anticancer abilities of *TIMP-3*.

#### Cell proliferation and apoptosis

Cancer cells grow in an uncontrolled fashion and evade the host immune system. Overexpression of *TIMP-3* suppresses proliferation and induces apoptosis in different cancer cell lines.<sup>62</sup> *TIMP-3* has been shown to induce cell apoptosis by activating mitochondrion-mediated caspase-3 in highly metastatic prostate cancer cell lines PC-3 and DU-145.<sup>108</sup> In addition, *TIMP-3* can enhance the sensitivity of cancer cells to apoptosis by stabilizing the death receptor. Mark *et al.* reported that *TIMP-3* induces a type II apoptotic

**Table 1.** MicroRNAs target TIMP-3 during cancer progression.

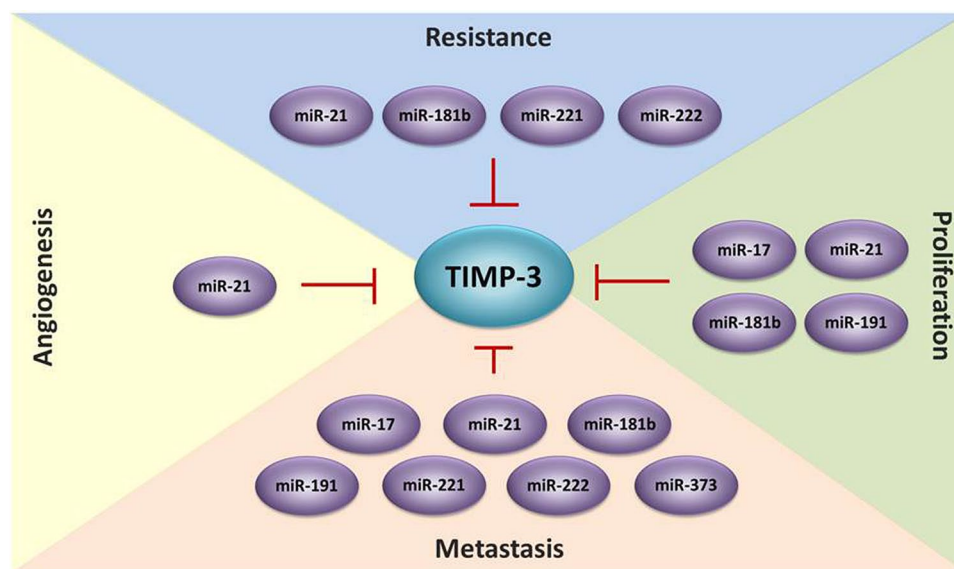
MicroRNAs	Cancer type	Function	Reference
miR-17-3p and miR-17-5p	Prostate cancer	Tumor growth and invasion ↑	Yang <i>et al.</i> <sup>85</sup>
miR-21	Melanoma	Tumor invasion ↑	Martin del Campo <i>et al.</i> <sup>86</sup>
	Breast cancer	Cancer metastasis ↑	Li <i>et al.</i> <sup>87</sup>
	Cervical cancer ESCC PDAC	Proliferation, migration and invasion ↑ Tumor growth and invasion ↑ Apoptosis ↓ Worse survival of patients	Zhang <i>et al.</i> <sup>88</sup> Wang <i>et al.</i> <sup>89</sup> Nagao <i>et al.</i> <sup>91</sup>
miR-21-5p	Gastric cancer	Doxorubicin resistance	Chen <i>et al.</i> <sup>92</sup>
	Breast cancer	Tumor growth ↑ Angiogenesis ↑	Dai <i>et al.</i> <sup>93</sup>
miR-181b	HCC	Tumor growth, migration and invasion ↑ Doxorubicin resistance	Wang <i>et al.</i> <sup>94</sup>
	Gastric cancer	Cancer metastasis ↑	Zhou <i>et al.</i> <sup>95</sup>
miR-191	EAO	Proliferation and invasion ↑	Dong <i>et al.</i> <sup>96</sup>
miR-221	OSCC	Doxorubicin resistance	Du <i>et al.</i> <sup>97</sup>
		Adriamycin resistance	Chen <i>et al.</i> <sup>98</sup>
miR-221 and miR-222	NSCLC and HCC	TRAIL resistance	Garofalo <i>et al.</i> <sup>99</sup>
	Breast cancer	Migration ↑ Tamoxifen resistance	Gan <i>et al.</i> <sup>100</sup>
miR-373	ESCC	Migration and invasion ↑	Liu <i>et al.</i> <sup>101</sup>

EAO, endometriosis-associated ovarian cancer; ESCC, esophageal squamous cell carcinoma; HCC, hepatocellular carcinoma; NSCLC, non-small cell lung cancer; OSCC, oral squamous cell carcinoma; PDAC, pancreatic ductal adenocarcinoma; TRAIL, TNF-related apoptosis-inducing ligand.

pathway *via* a Fas-associated death domain-mediated mechanism.<sup>109</sup> TIMP-3 overexpression in melanoma cells induces apoptosis by stabilizing the TNF receptor-1 (R1), FAS, and TNF-related apoptosis-inducing ligand receptor-1 (TRAIL-R1) on the cell surface; stabilization of death receptors results in the activation of apoptosis markers caspase-8 and caspase-3.<sup>110</sup> Other studies have also suggested that TIMP-3 can block ADAM17 activity, which induces shedding of TNF-R1 and TNF- $\alpha$  from the cell surface.<sup>111,112</sup> Notably, TIMP-3 can also promote death of nonadherent small-cell lung carcinoma cells even if the cells do not present cell surface death receptors or caspase-8.<sup>113</sup> *In vivo* studies have shown that prostate cancer cells transfected with TIMP-3 suppress tumor growth and induce tumor apoptosis in nude mice.<sup>114</sup> Similar results have also been shown in colon carcinoma, melanoma, and neuroblastoma; TIMP-3 overexpression suppresses tumor growth *in vivo*.<sup>15,115,116</sup>

### Angiogenesis

Angiogenesis, which is the process of new blood vessel formation, plays a key role in tumor growth and metastasis. TIMP-3 has been shown to inhibit angiogenesis through regulating angiogenesis-related proteins or directly inducing apoptosis of endothelial cells. Vascular endothelial growth factor (VEGF) is a key mediator of blood vessel development. TIMP-3 can inhibit angiogenesis by blocking VEGF from binding to VEGF receptor-2.<sup>117</sup> In leukemia cells, TIMP-3 can inhibit the proliferation and migration of human umbilical vein endothelial cells (HUVECs) and reduce VEGF-mediated MMP-2 and MMP-9 expression.<sup>67</sup> Kang *et al.* used a yeast 2-hybrid system and found a TIMP-3-interacting partner angiotensin II type 2 receptor (AGTR2). They suggested that combination treatment with TIMP-3 and AGTR2 inhibits VEGF-mediated proliferation of HUVECs.<sup>118</sup> TIMP-3 can also inhibit angiogenesis by promoting endothelial cell



**Figure 1. TIMP-3 is regulated by microRNAs during cancer progression.** MicroRNAs play an important role to silence the expression of TIMP-3 and promote cell proliferation, migration, invasion, metastasis, angiogenesis and drug resistance in several types of cancer.

apoptosis. Qi *et al.* suggested that TIMP-3 may induce apoptosis of endothelial cells by triggering a FAK/Paxillin cell survival pathway but not a caspase-dependent cell death pathway.<sup>119</sup> Upregulation of TIMP-3 by transcription factor sp1 *via* P14ARF inhibited endothelial cell migration and vessel formation.<sup>34</sup> In an *in vivo* study, *TIMP-3* knockout in nude mice but not in the tumor showed enhanced growth of tumor and increased angiogenesis.<sup>120</sup>

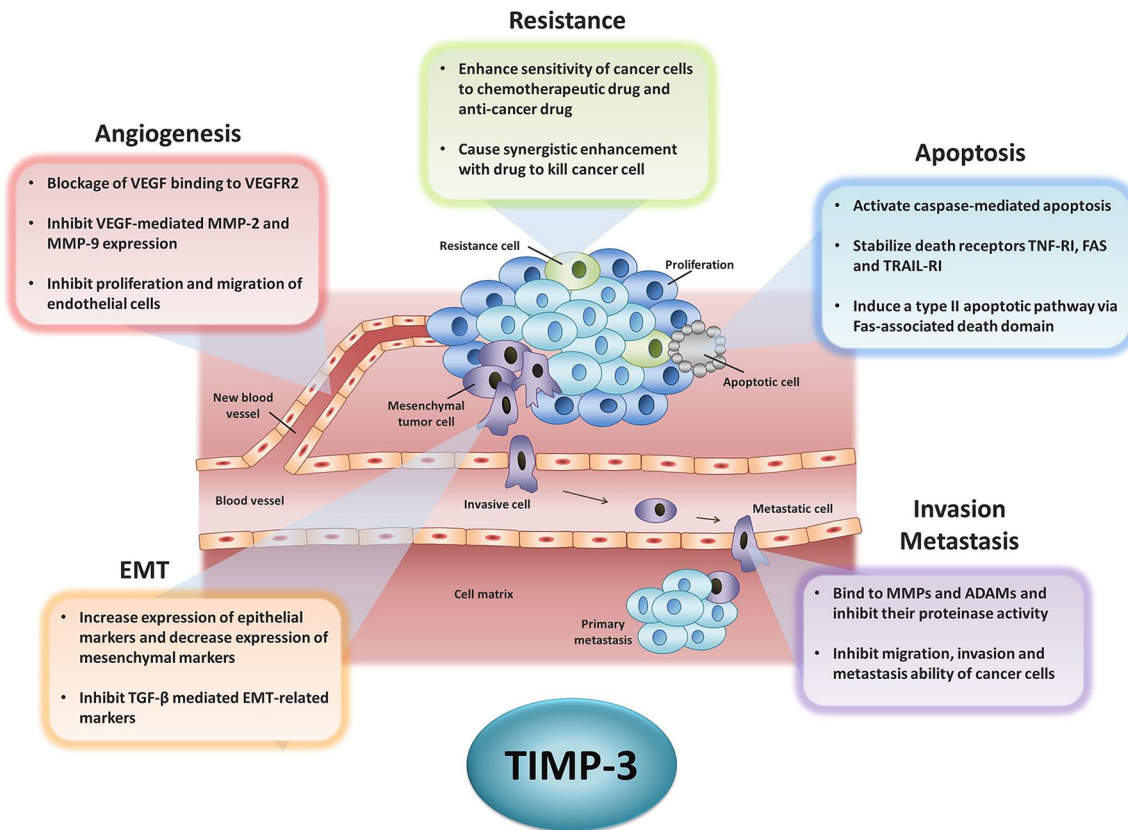
#### Migration and invasion and metastasis

Metastasis is the process of cancer cells breaking apart from the primary tumor and traveling through the blood or lymph vessels to form new tumors in other organs or tissues.<sup>2,7,121,122</sup> Metastasizing cancer cells can secrete proteinases to degrade the ECM and further migrate and invade into the blood. Overexpression and unrestricted activation of MMPs may promote malignant conversion of cancer cells.<sup>123,124</sup> TIMP-3 has been shown to block proteinase activity and inhibit tumor migration and invasion. Overexpression of TIMP-3 in osteosarcoma decreases MMP-1 and MMP-2 expression and suppresses cell migration and invasion ability.<sup>66</sup> In addition, TIMP-3 can protect ECM degradation mediated by ADAM12, the expression of which is correlated with the status and stage of breast cancer.<sup>48</sup> Inhibition of the

invasive activity of melanoma cells is more pronounced in the case of TIMP-3 overexpression compared with TIMP-1 and TIMP-2 overexpression.<sup>16</sup> Loss of TIMP-3 promotes NSCLC cell invasion *via* TNF-mediated IL-6 production.<sup>24</sup> High levels of soluble CD44 are associated with malignant cancer and metastasis; however, TIMP-3 potentially inhibits CD44 shedding by targeting ADAM-like proteases and MT1-MMP.<sup>125,126</sup> In an animal study, melanoma and lymphoma cells in *TIMP-3*<sup>-/-</sup> mice were observed to have higher metastatic ability, metastasizing to multiple organs, and lung tissues from *TIMP-3*<sup>-/-</sup> mice showed higher MMP-2 and MMP-9 enzyme activity than did those from wide-type mice.<sup>127</sup>

#### EMT

EMT refers to the transition of an epithelial cell into a mesenchymal cell and is considered an important indicator of cancer cell metastasis.<sup>128,129</sup> Restoration of TIMP-3 expression in thyroid tumor cells increased their cell adhesion ability, thereby increasing the expression of the epithelial marker cytokeratin 8/18 and decreasing that of the mesenchymal marker vimentin.<sup>19</sup> In addition, transfection of pcDNA-*TIMP-3* in gastric cancer cells followed by treatment with TGF- $\beta$  revealed that TIMP-3 partially rescues EMT-related marker expression induced by TGF- $\beta$ .<sup>95</sup>



**Figure 2. The role of TIMP-3 in cancer progression.** Tumor development contains many complex mechanisms including avoiding from cell apoptosis, growth without limit, angiogenesis, resistance to drug treatment, changing cell morphology from epithelial type to mesenchymal type (EMT), and metastasizing to the new organ. This figure lists the anticancer capacity of TIMP-3 that has been reported in previous studies. ADAM, a disintegrin and metalloproteinase; EMT, epithelial–mesenchymal transition; MMP, matrix metalloproteinase; TGF, transforming growth factor; TIMP, tissue inhibitors of metalloproteinase; TNF, tumor necrosis factor; VEGF, vascular endothelial growth factor.

### TIMP-3 as a target for cancer therapy

In addition to traditional surgery, chemotherapy, and radiotherapy, current cancer research is increasingly focusing on identifying biomarkers for early prediction of cancer progression, immunotherapy, and the use of targeted therapy against tumor suppressor genes or oncogenes.

### *TIMP-3 as a biomarker for predicting cancer progression*

Cancer may rapidly progress from an early stage to an advanced stage, which hinders early diagnosis and accurate monitoring of progression. Plasma and serum levels of TIMP-3 can be easily detected using enzyme-linked immunosorbent assay. A clinical study showed that the increase in plasma levels of TIMP-3 was significantly higher in those with large tumors (>T2) than in those with small tumors among betel quid chewers with

oral cancer.<sup>23</sup> TIMP-3 protein and mRNA can be extracted from tissues of patient with cancer and detected using Western blotting, immunohistochemistry, and real-time polymerase chain reaction. TIMP-3 protein expression has been shown to be correlated with tumor stage in NSCLC.<sup>24</sup> Low TIMP-3 mRNA expressions was observed in patients with high-grade clear cell renal cell carcinoma.<sup>59</sup> Loss of TIMP-3 expression in gastric cancer tissues is associated with tumor size, histologic grade, lymphatic invasion, venous invasion, invasive depth, lymph node metastasis, distant metastasis, and TNM stage.<sup>25</sup> Gene polymorphism and DNA methylation could be identified from DNA extracted from blood, body fluids, and salivary rinse and used as biomarkers to predict cancer progression in many cancers. TIMP-3 polymorphism rs9862 has been associated with an increased risk of developing a tumor of size >T2 among betel quid chewers with oral

**Table 2.** TIMP-3 as biomarker in cancer progression.

Cancer type	Sample source	Clinical significance	Reference
<b>TIMP-3 expression</b>			
Oral cancer	Plasma protein	Tumor size	Su <i>et al.</i> <sup>21</sup>
NSCLC	Tissue protein	Tumor stage	Wu <i>et al.</i> <sup>24</sup>
CCRCC	Tissue mRNA	Tumor grade	Gu <i>et al.</i> <sup>58</sup>
Gastric cancer	Tissue protein	Tumor size Histologic grade Distant metastasis TNM stage	Yu <i>et al.</i> <sup>25</sup>
<b>TIMP-3 polymorphism</b>			
Oral cancer	Blood DNA	Tumor size	Su <i>et al.</i> <sup>23</sup>
Breast cancer	Blood DNA	Survival	Bashash <i>et al.</i> <sup>77</sup>
Adenocarcinoma	Blood DNA	Survival	Wieczorek <i>et al.</i> <sup>76</sup>
<b>TIMP-3 methylation</b>			
ESCC	Tissue DNA	Disease-free survival Overall survival	Ninomiya <i>et al.</i> <sup>130</sup>
AML	Bone marrow DNA	Cytogenetic prognosis	Raneros <i>et al.</i> <sup>131</sup>
HNSCC	Salivary rinse DNA	Local recurrence-free survival	Sun <i>et al.</i> <sup>132</sup>
Gastric cancer	Body fluid DNA	Disease-free survival Tumor size Differentiation T stage Lymph node metastasis Distant metastasis	Yu <i>et al.</i> <sup>25</sup>
AML, acute myeloid leukemia; CCRCC, clear cell renal cell carcinoma; ESCC, esophageal squamous cell carcinoma; HNSCC, head and neck squamous cell carcinoma; NSCLC, non-small cell lung cancer.			

cancer.<sup>23</sup> A study revealed that TIMP-3 hypermethylation in patients with esophageal squamous cell carcinoma is associated with poorer prognosis for both disease-free and overall survival than that of patients without TIMP-3 methylation.<sup>130</sup> Patients with AML harboring methylation of TIMP-3 show higher frequency of adverse cytogenetic prognosis than those with a favorable or intermediate prognosis.<sup>131</sup> TIMP-3 promoter hypermethylation detected from pre-treatment salivary rinse is significantly associated with local recurrence-free survival in patients with head and neck squamous cell carcinoma.<sup>132</sup> Methylation of the *TIMP-3* promoter identified from body fluids has been reported to be a useful biomarker for predicting tumor size, differentiation, T stage, lymph node metastasis, distant

metastasis, and clinical stage in patients with gastric cancer<sup>25</sup> (Table 2).

#### *Combination of TIMP-3 and clinical treatment in cancer therapy*

Cancers cells can develop drug resistance, which could limit the efficacy of traditional therapies. Combination therapy, the use of more than one type of therapy in treatment, has the potential to delay or reduce drug resistance of cancer. In prostate cancer, adenovirus-mediated expression of TIMP-3 highly sensitizes prostate cancer cells to the chemotherapeutic drug paclitaxel (Taxol) and promotes synergistic enhancement in cell death.<sup>108</sup> TIMP-3 increases cisplatin-induced apoptosis in laryngeal carcinoma by facilitating a



mitochondria-dependent apoptosis mechanism such as cytochrome c release and caspase activation.<sup>133</sup> In addition, TIMP-3 overexpression in osteosarcoma cells facilitates cisplatin-induced apoptosis, whereas TIMP-3 knockdown by siRNA has an opposite effect.<sup>66</sup> In an animal study, combination therapy with adenovirus-TIMP-3 and the broad-spectrum antitumor agent cisplatin inhibited cervical cancer xenograft growth more effectively than cisplatin only.<sup>134</sup>

#### *Inhibition of cancer progression through TIMP-3*

Because TIMP-3 expression is downregulated or deregulated in cancer, current studies are focusing on normalizing or reactivating the expression of TIMP-3 (Table 3). In recent years, the use of natural products has gained much attention in cancer chemoprevention. For example, Andrographolide, a Chinese herbal medicine that is isolated from the stem and leaves of *Andrographis paniculata*, can inhibit angiogenesis

by suppressing miR-21-5p expression and enhancing TIMP-3 expression.<sup>93</sup> Diallyl disulfide, one of the organosulfur compounds derived from *Allium* vegetables, can inhibit migration and invasion in gastric cancer and upregulate tumor suppressor gene expression, including that of TIMP-3 and E-cadherin.<sup>135</sup> Green tea polyphenols and their major component epigallocatechin-3-gallate restore TIMP-3 expression by attenuating epigenetic silencing of EZH2 and HDACs, thus inhibiting invasion in breast cancer.<sup>136</sup> Arctigenin, derived from the seeds of *Arctium lappa*, may also increase TIMP-3 expression and inhibit tumor growth in prostate cancer.<sup>137</sup> Mithramycin A (MMA), an antibiotic against Gram-positive soil bacteria, has antitumor and antimetastatic effects; MMA can interact with the catalytic pocket of DNMT1, which results in promoter demethylation and mRNA restoration of TIMP-3.<sup>138</sup> Although beneficial, cancer chemoprevention is associated with many challenges, such as toxicity, side effects, efficacy only at high doses, low

**Table 3.** Anticancer agents targeting to TIMP-3.

Agent name	Targeting cancer	Anticancer function	Reference
<b>Natural products</b>			
Andrographolide	Breast cancer	Antiangiogenesis	Dai <i>et al.</i> <sup>93</sup>
Diallyl disulfide	Gastric cancer	Inhibit migration and invasion	Su <i>et al.</i> <sup>135</sup>
EGCG	Breast cancer	Inhibit invasion	Deb <i>et al.</i> <sup>136</sup>
Arctigenin	Prostate cancer	Inhibit tumor growth	Wang <i>et al.</i> <sup>137</sup>
Mithramycin A	Lung cancer	Inhibit metastasis	Lin <i>et al.</i> <sup>138</sup>
<b>Synthetic products</b>			
NucAnt 6L	Melanoma	Inhibit invasion	Destouches <i>et al.</i> <sup>140</sup>
p700	Breast cancer	Inhibit tumor growth and angiogenesis	Chen <i>et al.</i> <sup>141</sup>
lncRNA	Diffuse large B-cell lymphoma	Induce apoptosis and inhibit tumor growth	Su <i>et al.</i> <sup>142</sup>
TAPI-0	AML	Inhibit AML cell-induced STNK cell abnormalities	Arriga <i>et al.</i> <sup>143</sup>
<b>Epigenetic agents</b>			
SGI-1027	Multiple cancers	Block DNMT1 and reactivate TIMP-3	Datta <i>et al.</i> <sup>144</sup>
MPT0G013	Colon cancer	Inhibit angiogenesis, tumor growth, metastasis	Wang <i>et al.</i> <sup>145</sup>
decitabine	AML	Enhance the lytic activity of NK cells	Raneros <i>et al.</i> <sup>131</sup>
DNMT1, DNA methyltransferase 1; EGCG, epigallocatechin-3-gallate; NK, natural killer; p700, synthetic peptide derived from N-terminal domain of TIMP-3; STNK, short-term natural killer.			

bioavailability, and rapid metabolism.<sup>139</sup> In addition to the natural products, some synthetic products can also be used against cancer. NucAnt 6L (N6L), a synthetic peptide, can suppress the invasion ability of melanoma through TIMP-3 release from sulfated glycosaminoglycans.<sup>140</sup> The synthetic peptide p700, which is derived from the N-terminal domain of TIMP-3, inhibits VEGF-family receptors, angiogenesis, and tumor growth.<sup>141</sup> An artificially designed i-lncRNA can target multiple oncogenic miRNAs, thereby protecting the tumor suppressor gene TIMP-3 in diffuse large B-cell lymphoma.<sup>142</sup> TAPI-0, a functional analog of TIMP-3, which inhibits AML cell-induced short-term natural killer (NK) cell abnormalities, may contribute to NK cell-based immunotherapy of AML.<sup>143</sup> Other studies have focused on the epigenetic regulation of TIMP-3. Jharna *et al.* suggest that the quinoline-based compound SGI-1027 acts as a hypomethylation agent by inhibiting DNMT1 activity and inducing its degradation. In addition, they claimed that treatment with SGI-1027 facilitates re-expression of the tumor suppressor genes MLH1, P16, and TIMP-3 in cancer cell lines.<sup>144</sup> MPT0G013, a HDAC inhibitor, has been reported to induce TIMP-3 expression and further inhibit angiogenesis and tumor growth *in vivo*.<sup>145</sup> Moreover, TIMP-3 may be a target for demethylation treatments in AML patients. TIMP-3 demethylation by azacytidine (AZA) or decitabine (DAC) can inhibit the release of soluble NKG2D ligand (NKG2DL) mediated by ADAM17 and enhance the lytic activity of NK cells through immune recognition mediated by the NKG2D–NKG2DL engagement.<sup>131</sup> Although hypomethylating agents (HMAs) such as AZA and DAC have been widely used in the clinical treatment of AML. However, there are still many limitations and side effects in the use of HMAs,<sup>146</sup> and it is easy to cause excessive inhibition of DNA methylation. Moreover, DNA hypomethylation may cause some autoimmune diseases such as systemic lupus erythematosus (SLE),<sup>147,148</sup> and may also activate some cancer metastasis genes such as u-PA to accelerate cancer metastasis.<sup>149</sup> Therefore, second-generation HMAs and other combined treatments are still evolving and ultimately expected to improve efficacy and reduce side effects.<sup>150</sup>

### Conclusion

TIMP-3 is unique among the TIMP family members because it is the only TIMP that can bind firmly to the ECM after secretion. In addition,

TIMP-3 can inhibit not only MMPs but also a wide range of ADAMs and ADAMTSs. Accumulating evidence indicates that the *TIMP-3* gene acts as a tumor suppressor gene by inducing apoptosis and inhibiting proliferation, angiogenesis, and metastasis. However, TIMP-3 expression is downregulated by genetic and epigenetic alternation in most cancers. In this review, we have systematically described the contribution of TIMP-3 in cancer and its potential in cancer therapy including its utility as a predictor of cancer progression, its efficacy in combination therapy for cancer treatment, and its potential as a direct target for cancer therapy. Further research is required to harness the potential of TIMP-3 in diagnosing and treating cancer.

### Funding

The author(s) received no financial support for the research, authorship, and/or publication of this article.

### Conflict of interest statement

The authors declare that there is no conflict of interest.

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