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## Biochemical role of the collagen-rich tumour microenvironment in pancreatic cancer progression

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### Abstract

PDAC (pancreatic ductal adenocarcinoma) is among the most deadly of human malignancies. A hallmark of the disease is a pronounced collagen-rich fibrotic extracellular matrix known as the desmoplastic reaction. Intriguingly, it is precisely these areas of fibrosis in which human PDAC tumours demonstrate increased expression of a key collagenase, MT1-MMP [membrane-type 1 MMP (matrix metalloproteinase); also known as MMP-14]. Furthermore, a cytokine known to mediate fibrosis *in vivo*, TGF- $\beta$ 1 (transforming growth factor- $\beta$ 1), is up-regulated in human PDAC tumours and can promote MT1-MMP expression. In the present review, we examine the regulation of PDAC progression through the interplay between type I collagen (the most common extracellular matrix present in human PDAC tumours), MT1-MMP and TGF- $\beta$ 1. Specifically, we examine the way in which signalling events through these pathways mediates invasion, regulates microRNAs and contributes to chemoresistance.

### Keywords

chemoresistance; fibrosis; microRNA; membrane-type 1 matrix metalloproteinase (MT1-MMP); pancreatic cancer; Snail; transforming growth factor- $\beta$  (TGF- $\beta$ )

## INTRODUCTION

PDAC (pancreatic ductal adenocarcinoma) is among the most aggressive of human malignancies and a leading cause of cancer-related mortality. Worldwide, there are over 200000 deaths from PDAC each year, with 38000 annual deaths in the U.S.A. alone, as well as over 60000 deaths in Europe [1-3]. Yet what is perhaps most striking about this disease, among other cancers with a dismal prognosis, is the rapidity of its inexorable progression

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[4,5]. Despite the use of intense chemotherapy and targeted therapies, the median survival of patients with PDAC is less than 1 year. Overall 5-year survival is less than 5% [4,5]. Over 80% of patients with PDAC present with disease that is not amenable to surgical resection because of either advanced local tumour growth or spread to distant sites. Even for those who do undergo surgery, treatment remains arduous with a 5-year survival of only 20% [6]. Several factors are thought to contribute to the aggressive nature of PDAC. The location of the pancreas deep within the abdomen means patients are often asymptomatic until the disease is too advanced for curative options. Unlike some malignancies in which patients present on screening examinations with small tumours that can be easily excised without harming surrounding tissue, many pancreatic cancer patients come to medical attention because of jaundice from obstruction of the bile ducts or pain from invasion of the surrounding nerves [4,5]. By this stage, the tumour has typically infiltrated the pancreaticobiliary system, often encasing vital blood vessels in the vicinity, thereby excluding surgical resection. In addition, in a unique molecular hallmark, PDAC is also known to be associated with a dense collagen-rich fibrotic reaction (Figure 1) [7-9]. This so-called desmoplastic reaction is known to contribute to disease progression and chemoresistance [10,11].

## ROLE OF THE DESMOPLASTIC REACTION IN PANCREATIC CANCER

Human PDAC tumours are predominantly composed of non-neoplastic inflammatory and fibroblastic cells and contain tremendous quantities of collagen-rich ECM (extracellular matrix) [7,8]. Unlike most other tumours, this desmoplastic reaction is a characteristic feature of human PDAC and is most striking when seen side-by-side with areas of relatively normal pancreas in which only minimal fibrosis is observed [7,8]. Importantly, transgenic mouse models have been able to recapitulate the stromal changes seen in human PDAC tumours. For example, PDAC tumours developing in mice expressing activated K-Ras<sup>G12D</sup> together with either inactivation of the Ink-4a/Arf tumour suppressor locus or p53 demonstrate significant desmoplasia with increased collagen expression [12-15]. In human disease, the correlate for this finding is an increased amount of interstitial fibrillar collagen (types I and III) as compared with normal pancreas [16,17]. Furthermore, during PDAC progression the normal basement membrane architecture is lost and malignant cells are directly exposed to interstitial collagen [16-18].

### Myofibroblasts and the desmoplastic reaction

Beyond changes in the composition of the ECM, the desmoplastic reaction is also associated with accelerated proliferation of fibroblasts, in some cases outnumbering local tumour cells [19,20]. A more detailed molecular analysis demonstrates that these proliferating cells are mesenchymal (or stellate) cells that have differentiated into an activated myofibroblastic phenotype [19-21]. As activated myofibroblasts, they exhibit increased expression of the cytoskeletal protein  $\alpha$ -smooth muscle actin and have been identified as the primary source of type I collagen that is present in the desmoplastic reaction. Importantly, there is significant cross-talk between fibroblasts and cancer cells, as the invasive potential of PDAC cells is greatly enhanced by coculture with stromal fibroblasts [22,23]. Co-injection of PDAC tumour cells with human stellate cells in an orthotopic mouse model results in

increased primary tumour incidence, size and metastasis, notably with a direct correlation with the proportion of stellate cells present [23]. Genes encoding for proteins already known to be associated with tumour invasion, metastasis and angiogenesis are also differentially expressed in both PDAC cells and fibroblasts as a consequence of this interaction [24].

### Fibrillar collagen and PDAC progression

PDAC is a malignancy with a widely recognized metastatic potential, yet one of the hallmarks of the disease is a dense fibrotic ECM that would seem to serve as a barrier to invasion. Indeed, interstitial collagens are highly resistant to proteolysis owing to their triple helical structure and fibrillar organization [25-27]. Breakdown of fibrillar collagen helices requires the enzymatic activity of specific collagenases, including interstitial collagenase [MMP (matrix metalloproteinase)-1], neutrophil collagenase (MMP-8), collagenase 3 (MMP-13) and MT1 (membrane type 1)-MMP [28,29]. Cathepsins have also been implicated in collagen cleavage, particularly under conditions of acidic pH [30]. On the basis of structural assessment alone, it would seem that the desmoplastic reaction should be a barrier and not an aid to PDAC progression. However, in a paradoxical twist, multiple studies have shown that type I collagen enhances tumour progression. Instead of blocking tumour cell motility, it is now clear that collagen fibres can act as highways for migration and facilitate metastasis by directing cancer cells to the all-important vasculature needed for local proliferation and distant spread [31,32]. In keeping with these findings, increased expression of type I collagen is detected in gene expression signatures associated with increased risk of metastasis [33-35]. Our own work has also shown that type I collagen can affect PDAC behaviour by increasing cellular motility and promoting expression of MT1-MMP [36,37].

It is also possible that the desmoplastic reaction in pancreatic tumours contributes to increased tumour migration and invasion through increased expression of an MMP-resistant isoform of type I collagen [38,39]. The normal isoform of type I collagen is an  $\alpha 1_2\alpha 2$  heterotrimer consisting of two  $\alpha 1(I)$  chains and one  $\alpha 2(I)$  chain [25,40]. However, an  $\alpha 1_3$  homotrimeric isoform of type I collagen composed of three  $\alpha 1(I)$  chains has been found in carcinomas *in vivo* and in cancer cells grown *in vitro* [41-46]. Although it remains unknown whether or not human PDACs express the  $\alpha 1_3$  homotrimeric collagen isoform, these homotrimers are resistant to all collagenolytic MMPs and to degradation by cancer cells and fibroblasts [38,39]. Importantly, cancer cells demonstrate enhanced migration atop such atypical homotrimeric type I collagen fibres when compared with heterotrimeric type I collagen fibres found in normal tissue [38,47]. However, in the multicellular milieu of a developing carcinoma, collagen homotrimers are produced primarily by cancer cells and not by cancer-associated fibroblasts [38], which primarily contribute to the stromal reaction in pancreatic cancer [8,48]. As a result, collagen homotrimers probably comprise only a small, yet significant, pro-tumorigenic fraction of the different collagens found in the heterogeneous environment of tumour tissue.

Type I collagen can also regulate cellular proliferation by modulating intracellular signals transduced by cell-surface receptors [49]. Notably, upon binding to integrins or discoidin domain receptors, type I collagen activates a series of intracellular signalling events with

profound effects on cell-cycle regulatory proteins, cyclins and CDKs (cyclin-dependent kinases) [50,51]. The activation of these signalling pathways is highly dependent upon the physical state of type I collagen: fibrillar collagen may inhibit cell growth, whereas non-fibrillar collagen may have no effect. More specifically, fibrillar type I collagen inhibits vascular smooth muscle cell proliferation by increasing the levels of the CDK inhibitors p21<sup>Cip1</sup> and p27<sup>Kip1</sup> [52]. Moreover, type I collagen can induce opposing cellular effects based upon the tensile strength and structure of the extracellular milieu [53]. When grown in two-dimensional collagen films, hepatocytes demonstrate enhanced cyclin D1 expression and DNA synthesis, yet, when grown in three-dimensional collagen gels, cyclin D1 expression and DNA synthesis are inhibited [54]. This difference in cyclin D1 expression between the two-dimensional and three-dimensional microenvironments was shown to result from differential activation of MAPK (mitogen-activated protein kinase) signalling. Hepatocytes on two-dimensional collagen films demonstrate increased ERK (extracellular-signal-regulated kinase) 1/2 phosphorylation compared with cells in three-dimensional collagen gel; inhibiting ERK1/2 activation with PD98059 blocks both cyclin D1 expression and DNA synthesis [54]. Furthermore, as the structural rigidity of the ECM increases, integrin-mediated intracellular signalling can be induced [55-57]. As the ECM becomes less flexible, force-dependent integrin clustering in breast cancer cells increases, in turn enhancing ERK1/2 activation and cell motility [55,56]. Since the desmoplastic reaction in pancreatic tumours is associated with increased cross-linking of fibrillar collagen (and thereby increased rigidity of the ECM), the collagen-rich desmoplastic reaction may also contribute to PDAC progression by enhancing tension-dependent integrin-mediated signalling in PDAC cells.

### The desmoplastic reaction and chemoresistance

The fibrotic extracellular environment that develops in the setting of PDAC does not appear to prevent tumour cell invasion out of the site of the primary tumour, but more and more data have now implicated the desmoplastic reaction as a substantial barrier to the effective targeting of chemotherapeutic agents into the site of primary disease. In a transgenic mouse model of PDAC, both delivery and efficacy of gemcitabine were improved following treatment with Hedgehog signalling pathway inhibitors which deplete the tumour-associated stromal reaction [10]. We have found that pancreatic cancer cells grown in three-dimensional collagen demonstrate decreased sensitivity to gemcitabine chemotherapy and continue to proliferate despite drug treatment [11]. In contrast with the effects of three-dimensional collagen on ERK1/2 signalling in hepatocytes, PDAC cells embedded in three-dimensional collagen show increased ERK1/2 signalling and subsequent attenuation of the effects of gemcitabine. Inhibiting ERK1/2 phosphorylation sensitizes PDAC cells in three-dimensional collagen to gemcitabine and causes growth arrest. ERK1/2 signalling is known to promote chemotherapy resistance in several different cancers, including multiple myeloma [58], HCC (hepatocellular carcinoma) [59] and PDAC [60]. Interestingly, in PDAC models, the prochemoresistance effect of collagen is only seen when the cells are grown in three-dimensional collagen and not atop three-dimensional collagen gels [11]. We have found that this is not due to differences in drug delivery to cancer cells, but rather to the increased DNA damage response induced in the three-dimensional collagen microenvironment (S. Dangi-Garimella and H. G. Munshi, unpublished work). This finding

of type I collagen directly protecting cancer cells against chemotherapy has also been noted in other malignancies; previous studies in lung cancer models have shown that collagen provides survival signals to attenuate the effects of chemotherapy and to allow for continued tumour cell proliferation [61].

## ROLE OF MT1-MMP IN PDAC PROGRESSION

Results of gene expression studies of stromal and neoplastic cells at the site of primary PDAC invasion have shown that MT1-MMP, a primary interstitial collagenase, is overexpressed by pancreatic cancer cells [62]. Immunohistochemistry and *in situ* hybridization studies have further clarified that MT1-MMP is overexpressed in pancreatic tumours relative to normal pancreas and that expression of MT1-MMP is enhanced in metastatic PDAC lesions as compared with their primary tumour of origin [36,63-65]. Genetic studies support the role of MT1-MMP as a primary regulator of interstitial collagenolysis, as mice genetically deficient in MT1-MMP have severe growth defects due to their subsequent inability to process interstitial collagens during critical stages of bone and soft tissue formation [66,67]. Moreover, *in vitro* studies using organotypic and three-dimensional culture systems demonstrate that MT1-MMP is the only collagenase to confer a three-dimensional growth advantage, presumably via removing the structural confines of the ECM and enabling the structural changes in the geometry of the cellular microenvironment necessary to drive proliferative responses [68,69]. Interestingly, in these studies cells failed to proliferate when embedded in collagenase-resistant three-dimensional type I collagen gels, suggesting that cleavage of type I collagen by MT1-MMP is necessary for cellular proliferation in a collagen-dominated microenvironment [68]. Indeed, overexpression of MT1-MMP alone may be sufficient to induce tumour formation. Overexpression of MT1-MMP in mouse mammary gland using a transgenic model induced adenocarcinoma of the breast [70]. Similarly, overexpression of MT1-MMP in non-tumorigenic well-differentiated MDCK (Madin–Darby canine kidney) cells was sufficient to drive formation of invasive tumours in nude mice [71].

Beyond its role in facilitating growth and invasion in the collagen microenvironment, MT1-MMP has also been implicated in the generation of the collagen-rich stromal reaction. It was previously noted that expression of MT1-MMP in the mouse mammary gland led to significant fibrosis [70], whereas expression of MT1-MMP in MDCK, squamous cell carcinoma and pancreatic cancer cells caused significant fibrosis when the cells were injected *in vivo* [11,70-72]. Our own work has shown that expression of MT1-MMP in the mouse pancreas causes large lesions associated with a pronounced stromal reaction [73]. These lesions are further associated with an increased number of  $\alpha$ -smooth muscle actin(+) cells, findings consistent with the increased numbers of myofibroblasts found in human pancreatic tumours. The increased fibrosis is also associated with increased Smad2 phosphorylation [73], indicating increased TGF- $\beta$  (transforming growth factor  $\beta$ ) signalling in MT1-MMP-expressing mice. We have also shown that the positive association between MT1-MMP, fibrosis and increased Smad2 phosphorylation is present in human pancreatic cancer specimens [36], further suggesting significant cross-talk between MT1-MMP and TGF- $\beta$  signalling during PDAC progression (also see below).

## MT1-MMP and chemoresistance

In the complex interplay between the extracellular microenvironment and the proliferation and metastatic potential of pancreatic cancer cells, MT1-MMP can function to regulate the nuances of chemotherapy resistance generated by the collagen microenvironment. We recently showed that blocking MT1-MMP expression or function in three-dimensional collagen sensitizes PDAC cells to gemcitabine [11]. Although MT1-MMP does not affect the response to gemcitabine when cells are grown on two-dimensional surfaces, modulation of MT1-MMP does significantly affect the chemotherapy response in cells grown in three-dimensional collagen. It is now well established that intercellular focal adhesions in the three-dimensional setting have a different molecular composition when compared with intercellular two-dimensional focal adhesions, further underscoring the role of microenvironment-dependent signalling in the global context of cellular behaviour [74]. The functional implications of the extracellular milieu for cancer cell signalling and behaviour are only recently becoming more notable. In breast cancer cells, MT1-MMP can increase the functionality of the collagen-binding  $\alpha 2\beta 1$  integrin [75]. Additionally, the ILK (integrin-linked kinase), a kinase capable of interaction with the cytoplasmic tail of  $\beta 1$  integrin, has been shown to contribute to gemcitabine resistance in PDAC cells [76]. Taken together, these findings suggest that it may be possible that MT1-MMP potentiates integrin signalling only in the three-dimensional microenvironment, thus enhancing PDAC gemcitabine resistance.

Mechanistically, it has been established that MT1-MMP in the collagen microenvironment increases ERK1/2 signalling to promote gemcitabine resistance. Overexpression of MT1-MMP increases ERK1/2 phosphorylation and increases gemcitabine resistance, whereas siRNA (small interfering RNA) against MT1-MMP decreases ERK1/2 phosphorylation and sensitizes pancreatic cancer cells to gemcitabine [11]. Despite the fact that MT1-MMP has been reported to increase ERK1/2 signalling via the cytoplasmic tail [77], pancreatic cancer cells expressing MT1-MMP devoid of the cytoplasmic tail nonetheless demonstrate robust ERK1/2 phosphorylation in the collagen microenvironment [11]. Although it has also been reported that TIMP (tissue inhibitor of metalloproteinases)-2 binding to MT1-MMP induces ERK1/2 phosphorylation in the absence of MT1-MMP proteolytic activity [78], MT1-MMP-induced ERK1/2 phosphorylation in the collagen microenvironment does require the catalytic activity of this proteinase. MT1-MMP enhances ERK1/2 phosphorylation in HSC-4 oral cancer cells grown in three-dimensional collagen, with the increased ERK1/2 phosphorylation blocked by MMP inhibitors [79]. Increased ERK1/2 phosphorylation in three-dimensional collagen can also be blocked by both an  $\alpha v\beta 3$  function-blocking antibody and siRNAs directed against Src and paxillin [79], suggesting that MT1-MMP in three-dimensional collagen enhances integrin signalling. Additionally, we have found that the increased ERK1/2 phosphorylation in three-dimensional collagen can be attenuated by blocking TGF- $\beta$  signalling [80], indicating that MT1-MMP may also modulate growth factor signalling in three-dimensional collagen to increase ERK1/2 activation. Consistent with this hypothesis, PDGF- $\beta$  (platelet-derived growth factor- $\beta$ -induced ERK1/2 phosphorylation in VSMCs (vascular smooth muscle cells) in the collagen microenvironment is blocked by TIMP-2 and is significantly attenuated in MT1-MMP-null VSMCs [81]. Interestingly, PDGF- $\beta$ -induced ERK1/2 phosphorylation in MT1-MMP-null



VSMCs can be restored by introducing either full-length MT1-MMP or MT1-MMP devoid of its cytoplasmic tail, but not by introducing the catalytically inactive form of MT1-MMP [81]. Taken together, these results suggest that MT1-MMP modulates growth factor and/or integrin signalling in the collagen microenvironment to enhance ERK1/2 phosphorylation, and thereby promote chemoresistance.

MT1-MMP also increases expression of another key protein associated with chemoresistance: the HMGA2 (high-mobility group A2) protein (Figure 2) [11]. HMGA2 is regulated by *let-7* family microRNAs (see below) and is a non-histone DNA-binding protein involved in chromatin remodelling and gene transcription [82,83]. HMGA proteins induce conformational changes in bound DNA and promote recruitment of regulators of transcription [83]. In addition to their role in chromatin remodelling, HMGA proteins have also been shown to be involved in chemoresistance, specifically as a determinant of chemoresistance in PDAC [84]. HMGA2 is part of the base end-joining repair machinery that removes small damaged bases from DNA, and has an apurinic/apyrimidinic lyase activity which can protect against the effects of chemotherapy agents designed to target tumour cell DNA [85]. HMGA2 can also increase expression of ATM (ataxia telangiectasia mutated), the main cellular sensor of genotoxic stress, and thereby increasing resistance to DNA-damaging agents [86]. In addition, through a mechanism that remains as yet unknown, HMGA2 can also maintain cancer stem cells in the undifferentiated state [82]. Stem cells are resistant to most cytotoxic chemotherapy [87], thus, in an example of the complexity of molecular circuitry, the extracellular collagenase MT1-MMP may ultimately play a role in inducing chemotherapy resistance in PDAC by increasing expression of an intracellular protein HMGA2 that subsequently modulates cancer stem cells.

### Regulation of MT1-MMP

MT1-MMP plays a critical physiological role as a major interstitial collagenase. Accordingly, its expression is highly regulated by several key transcription factors (Figure 3). Earlier reports have shown that culturing endothelial cells in three-dimensional collagen gels induced expression of Egr-1 leading to transcriptional activation of the MT1-MMP promoter, an association mediated in part by displacing the pre-bound basal transcription factor Sp1 [88,89]. Previously, it was demonstrated that three-dimensional collagen also induces Egr1 expression to subsequently promote MT1-MMP expression and invasion of ovarian cancer cells [90]. The three-dimensional collagen-induced Egr1 expression in ovarian cancer cells was mediated by  $\beta$ 1-integrin signalling through a Src kinase-dependent pathway [90]. MT1-MMP transcription can also be up-regulated by the well-known  $\beta$ -catenin–Tcf4 complex [91], placing MT1-MMP as a direct downstream target of the canonical Wnt signalling pathway. In addition, the transcription factor Snail, already known to be involved in the repression of E-cadherin during the pivotal EMT (epithelial–mesenchymal transition), can also up-regulate MT1-MMP expression [92,93]. Previous studies have shown that stable transfection of Snail in liver cancer cells increased MT1-MMP expression [92]. More recently, inducible expression of Snail in MCF7 breast cancer cells was also shown to increase MT1-MMP levels, thereby promoting growth and invasion in a chick chorioallantoic membrane assay [94]. Furthermore, fibroblasts in which Snail had been conditionally knocked out showed a decrease in MT1-MMP levels with attenuation of

invasion in the collagen microenvironment [95]. Our own studies have shown that Snail also increases MT1-MMP in pancreatic cancer cells, and blocking MT1-MMP with siRNA attenuates Snail-induced invasion [37]. Mechanistically Snail increases ERK1/2 phosphorylation, whereas blocking ERK1/2 activation inhibits both Snail-induced MT1-MMP expression and collagen invasion. Taken together, these findings demonstrate that crucial signalling pathways known to be important in tumour progression and invasion coordinately regulate MT1-MMP expression.

## ROLE OF TGF- $\beta$ IN THE DESMOPLASTIC REACTION

TGF- $\beta$  is frequently overexpressed in PDAC and is associated with both advanced tumour stage and a significant decrease in survival [8,96,97]. TGF- $\beta$  signals through cell-surface serine/threonine kinases to initiate cellular responses [98,99]. TGF- $\beta$  binding to its type II receptor (T $\beta$ RII) promotes T $\beta$ RII association with the type I receptor (T $\beta$ RI), followed by a phosphorylation event which then leads to subsequent phosphorylation of the R-Smads (receptor-associated Smads) Smad2 and Smad3. The phosphorylated R-Smads then bind to Smad4 and the complex translocates to the nucleus, wherein it can associate with transcription factors to regulate expression of target genes. Mutations in Smad4 are seen in an estimated 50% of PDAC tumours [100,101] and have been shown to affect tumour suppressive effects of TGF- $\beta$  signalling; restoration of Smad4 suppresses *in vivo* tumorigenicity [102]. Interestingly, it has been shown that, in a transgenic model of PDAC, selective deletion of Smad4 in the mouse pancreas alone has no effect on PDAC development [103]. Similarly, selective loss of T $\beta$ RII in the mouse pancreas also shows no discernible phenotype [104]. However, loss of Smad4 in combination with activated K-Ras<sup>G12D</sup> and other PDAC-relevant mutations (loss of Ink4a/Arf or p53) does result in rapid tumour development, providing genetic confirmation that Smad4 functions to block progression of K-Ras-initiated tumours [103,105]. In a similar fashion, loss of T $\beta$ RII in combination with activated K-Ras<sup>G12D</sup> also rapidly results in well-differentiated PDAC with 100% penetrance [104]. Interestingly, loss of Smad4 in the K-Ras<sup>G12D</sup>/Ink4a/Arf heterozygous mice not only leads to tumour development, but also results in an altered tumour phenotype, characterized by more well-differentiated tumours [103,105]. On the other hand, tumours with intact Smad4 show increased evidence of EMT, whereas a subset of advanced tumours with poor differentiation demonstrate TGF- $\beta$ -dependent growth [103,105]. Taken together, these findings indicate that TGF- $\beta$  and Smad4 may act further downstream as tumour promoters in the K-Ras<sup>G12D</sup>/Ink4a/Arf heterozygous mouse model.

In human PDAC specimens, there is a strong and well-known correlation between mRNA expression of TGF- $\beta$ 1 and type I collagen, suggesting that TGF- $\beta$ 1 may be directly responsible for the fibrotic reaction that develops in PDAC tumours [106,107]. Consistent with this hypothesis, TGF- $\beta$ 1 overexpression in PDAC cells induces the desmoplastic reaction in an experimental model of human pancreatic carcinoma [108]. Since fibroblasts are responsible for the synthesis, degradation and remodelling of much of the ECM, this finding indicates that TGF- $\beta$ 1 synthesized by pancreatic cancer can modulate fibroblast behaviour. Moreover, these fibroblasts can, in turn, modulate the behaviour of cancer cells



themselves through the secretion of additional cytokines and through further modification of the ECM microenvironment [109-115].

### **Cross-talk involving collagen, TGF- $\beta$ 1 and MT1-MMP**

Deciphering the nature of the complex signalling events resulting from cross-talk between components of the extracellular microenvironment, the proteinases capable of modulating this extracellular milieu and canonical cytokine signalling families well-known to regulate tumour cell growth and development is a challenging, but important, component of ongoing efforts to better understand the initiation and development of PDAC. Our own work has shown that human tumours with increased stromal reaction demonstrate increased TGF- $\beta$  signalling characterized by significant cross-talk between collagen, TGF- $\beta$ 1 and MT1-MMP [36]. Previously, we have published that pancreatic cancer cells induce TGF- $\beta$ 1 expression and signalling to promote MT1-MMP expression upon exposure to type I collagen [36]. Pancreatic cancer cells grown in three-dimensional collagen demonstrate increased TGF- $\beta$ 1 protein and mRNA levels, and increased Smad2 phosphorylation [36,80]. It was also previously shown that human mesangial cells cultured on type I collagen show increased TGF- $\beta$ 1 mRNA and protein levels [116]. Increased TGF- $\beta$ 1 production was diminished by a function-blocking anti- $\beta$ 1-integrin antibody and also by a dominant-negative ILK [116], indicating that type I collagen-induced TGF- $\beta$ 1 expression is regulated by  $\beta$ 1-integrins. However, beyond regulating TGF- $\beta$  expression, integrins can also promote activation of latent complexes of TGF- $\beta$  [117]. The  $\alpha$ v integrins, in particular  $\alpha$ v $\beta$ 3,  $\alpha$ v $\beta$ 5,  $\alpha$ v $\beta$ 6 and  $\alpha$ v $\beta$ 8, have been shown to liberate active TGF- $\beta$ 1 [118-121]. Significantly, mice carrying a single point mutation in the RGD integrin-binding motif of latent TGF- $\beta$ 1, thereby expressing latent TGF- $\beta$ 1 in a form that cannot be bound or activated by integrins, develop defects identical with those seen in mice completely lacking TGF- $\beta$ 1 [122]. Furthermore, generation of active TGF- $\beta$  in  $\alpha$ v $\beta$ 8-expressing cells requires proteolytic cleavage of LAP (latency-associated peptide) by MT1-MMP [121].

Our own work has also identified MT1-MMP/TGF- $\beta$ 1 cross-talk in a MT1-MMP transgenic mouse model. Expression of MT1-MMP in the mouse pancreas results in increased pancreatic fibrosis that is also associated with increased TGF- $\beta$  signalling [73]. Consistent with our observation, other groups have shown that expression of MT1-MMP in cardiac tissue causes significant fibrosis which is likewise associated with increased TGF- $\beta$  signalling [123]. Previous work has demonstrated that MT1-MMP can increase levels of the active-form of TGF- $\beta$  by cleaving latent TGF- $\beta$ -binding protein-1 to release TGF- $\beta$  bound to the ECM, an event which serves to increase the overall biological availability of TGF- $\beta$  [124-126]. As indicated above, MT1-MMP can also cleave the LAP itself to release the active-form of TGF- $\beta$  [121,123,126-128].

### **Cross-talk involving collagen, TGF- $\beta$ 1 and Snail**

Further complexity in the role of TGF- $\beta$  signalling in PDAC is introduced by considering the effects of this cytokine on the Snail family of transcription factors. TGF- $\beta$  is a well-defined inducer of Snail in a variety of cell lines, including human PDAC cells [129-131]. However, more recent studies have identified the additional mechanistic link: in a PDAC model system, collagen itself can activate TGF- $\beta$  signalling [36,37,80,116,132], in turn

leading to increased Snail expression (Figure 3) [37]. We have demonstrated that treatment with either a highly specific inhibitor of T $\beta$ RI or siRNA against T $\beta$ RI blocks collagen-induced Snail expression [37]. Moreover, knocking down Smad4 using siRNA also abrogates collagen-induced Snail expression in our model system. Interestingly, TGF- $\beta$ 1 can signal through both Smad2 and Smad3 to regulate Snail expression [131], but, in pancreatic cancer cells, collagen-induced Snail expression involves Smad3, but not Smad2 [37]. We have found that siRNA-mediated down-regulation of Smad3, but not Smad2, abrogates collagen-induced Snail expression [37]. A closer analysis of Smad2 and Smad3 reveals a differential role for these two transcription factors during both embryonic development and in functional adult tissue. Liver-specific knock down of Smad3 or Smad2 in adult mice has shown that Smad3, but not Smad2, is required for TGF- $\beta$ -stimulated EMT and growth arrest [133]. Similarly, TGF- $\beta$ 1-mediated repression of E-cadherin in kidney epithelial cells also requires Smad3, but not Smad2 [134,135]. On the basis of multiple studies in different tissue types, it appears that Smad3 may play a more significant role in regulating EMT than Smad2. However, other signalling events with the collagen microenvironment may also play a role in regulating this family of transcription factors, as collagen is also known to signal through ILK to regulate Snail expression [136-138]. Importantly, both integrins and integrin-associated proteins have been shown to regulate various growth factor signalling pathways including TGF- $\beta$  signalling [139,140]. The collagen integrin receptor  $\alpha$ 2 $\beta$ 1, known to associate with ILK [141], can also interact with the T $\beta$ RI-T $\beta$ RII heterodimeric complex to modulate Smad signalling [132,142].

### Snail regulation of fibrosis

Several studies in non-PDAC systems have implicated Snail as a regulator of extracellular fibrosis. Collagen is known to promote Snail expression, but fibrogenesis itself can also be initiated by Snail. CCl<sub>4</sub> (carbon tetrachloride)-induced fibrosis in the liver is associated with increased Snail expression, whereas liver-specific ablation of Snail attenuates this chemical-induced fibrosis [143]. Snail also promotes the activation of liver stellate cells, themselves important for activation of the fibrotic repair response during liver injury [144]. In adult renal cells, Snail activation is sufficient to drive both deposition of type I collagen and disruption of tissue architecture owing to the repression of cadherin-16 and the kidney differentiation factor HNF1 (hepatocyte nuclear factor 1) [145]. Given the clear role of Snail in regulating fibrosis in the kidney and liver, it is conceivable that Snail may also play an active role in signalling events driving the desmoplastic reaction found in PDAC tumours. In conjunction with the previously discussed cross-talk between Snail and other components of the extracellular PDAC milieu, this raises several intriguing questions for further study of the precise role played by this key transcription factor in the proliferative and metastatic events that characterize pancreatic malignancies.

### MicroRNA AND PANCREATIC CANCER

MicroRNAs are small, single-stranded, non-coding RNAs that can post-transcriptionally regulate gene expression [146,147]. The identification of microRNAs and subsequent efforts to understand their role in regulating the proliferation, differentiation and invasion events that characterize tumour cell behaviour represent a major focus of molecular biology

research during the last decade [148-151]. Not surprisingly, the role played by microRNAs in pancreatic cancer has been a rich area of investigation [152-155].

### Collagen regulation of microRNAs

We have previously shown that collagen represses the *let-7* family of microRNAs in pancreatic cancer cells. *Let-7* family members, which were initially identified as key regulators of embryonic development in *Caenorhabditis elegans* [156], can target HMGA2, K-Ras and number of other proteins [157-159]. They have also been shown to play an important role in human cancers. Expression of *let-7* increases with cell differentiation and is highest in mature tissues [160], in which microRNAs function to inhibit proliferation [161]. As with other key regulators of cell proliferation, aberrant expression of microRNA has been associated with human malignancies, including a noted decrease in the anti-proliferative *let-7* family in human PDAC samples [162]. Overexpression of *let-7* in mutant K-Ras-positive lung cancer cells inhibits tumour growth *in vivo* [163], and overexpression of *let-7* in breast tumour-initiating cells inhibits tumour formation and metastasis *in vivo* [164]. Interestingly, collagen-mediated repression of *let-7* involves MT1-MMP and TGF- $\beta$  signalling [80], both clearly established mediators of the pathogenesis of pancreatic malignancies. Our own work has shown that siRNA-mediated knock down of MT1-MMP attenuates collagen repression of *let-7* in PDAC cells [80]. Furthermore, blocking TGF- $\beta$  signalling similarly attenuates collagen-induced repression of *let-7*. Collagen repression of *let-7* in pancreatic cancer cells also involves ERK1/2 signalling [80], which our own work has shown is enhanced by MT1-MMP and TGF- $\beta$  signalling in the collagen microenvironment (Figure 2) [80]. Consistent with the report that ERK1/2 signalling regulates the human microRNA-generating complex to reduce *let-7* expression [165], collagen through increased ERK1/2 signalling also decreases *let-7* levels by inhibiting *let-7* processing to the mature form in pancreatic cancer cells [80]. These results further demonstrate the nuances of the significant cross-talk between collagen, MT1-MMP and TGF- $\beta$  signalling in the regulation of PDAC progression.

### MicroRNAs and fibrosis

MicroRNAs have also been implicated in the regulation of extracellular fibrosis, with several prominent examples standing out among previous studies (Table 1). First, *let-7d* expression is significantly decreased in lung samples from patients with IPF (idiopathic pulmonary fibrosis) [166]. *In vitro* inhibition of *let-7d* in lung epithelial cell lines increases the mesenchymal markers N-cadherin and vimentin, while decreasing expression of the epithelial marker cytokeratin [166]. Furthermore, inhibition of *let-7d in vivo* using antagomirs increases mesenchymal markers, decreases epithelial markers and increases collagen production, thus resulting in overall alveolar septal thickening. *Let-7g*, another member of the *let-7* family, is reduced in HCC and has been shown to inhibit expression of type I collagen [167]. Low levels of *let-7g* expression in HCC tumours were subsequently found to predict poor patient survival, and *in vitro* studies have shown that *let-7g* inhibits HCC cell growth and migration [167]. Expression of *miR-29*, an additional microRNA not associated with the *let-7* family, is reduced both in patients with advanced hepatic fibrosis and in a mouse model of CCl<sub>4</sub>-induced liver fibrosis [168]. This microRNA can also function to repress collagen production by stellate cells. Furthermore, overexpression of

*miR-29b* in immortalized murine hepatic stellate cells leads to a dose-dependent decrease in the expression of the collagen *Col1a1*, *Col4a5* and *Col5a3* genes [168]. In addition, in human stellate cells *miR-29b* was shown to be an effective suppressor of type I collagen at the mRNA level through direct binding to the *COL1A1* 3' UTR (untranslated region) [169]. Interestingly, Smad3 can also bind to the promoter region of *let-7d* to repress its expression [166], whereas TGF- $\beta$  represses *miR-29* expression [168].

The *miR-200* family of microRNAs also function to regulate fibrosis and have been identified as inhibitors of EMT in a number of different malignancies [170,171]. Recent work has shown that this family, in particular the *miR-200b* member, inhibits TGF- $\beta$ -induced EMT of renal tubular cells [172]. Intravenous injection of a *miR-200b* precursor in obstructed kidneys inhibits collagen I, collagen III and fibronectin, thereby reducing fibrosis [172]. Interestingly, the *miR-192* family member is a mediator of TGF- $\beta$ /Smad3-driven renal fibrosis through Smad3 binding to the *miR-192* promoter to increase collagen expression as a downstream effect [173]. Clearly, microRNAs share the same nuanced signalling events that characterize other families of intracellular and extracellular signalling modulators, even down to apparently paradoxical effects from very similar effectors.

### MicroRNAs and chemoresistance

Our own work has shown that *let-7* microRNA does not mediate gemcitabine resistance in pancreatic cancer either on two-dimensional surfaces or in the three-dimensional collagen microenvironment [80], but other microRNAs have been shown to mediate chemoresistance. For example, *miR-21*, overexpressed in PDAC tumours and a predictor of poor outcomes [174], contributes to gemcitabine resistance in PDAC, partially through modulation of PI3K (phosphoinositide 3-kinase)/Akt signalling [175,176]. *miR-21* increases pro-survival PI3K signalling through repression of PTEN (phosphatase and tensin homologue deleted on chromosome 10). It also up-regulates the pro-survival Bcl-2 protein and inhibits the pro-apoptotic Bax protein to promote gemcitabine resistance in PDAC cells. The effects of *miR-21* on PI3K/Akt and Bcl-2 signalling are not unique to pancreatic cancer cells, as *miR-21* can also mediate doxorubicin resistance in bladder cancer cells [177]. Interestingly, stem cells isolated from several different tumour types demonstrate up-regulation of both *miR-21* and its upstream regulator AP-1 (activator protein 1) [178]. Even more notable, a small-molecule inhibitor of AP-1 or an anti-microRNA against *miR-21* were both found to sensitize cancer stem cells to topotecan and decrease colony formation [178].

## CONCLUSIONS

The well-characterized desmoplastic reaction of pancreatic cancer has been described as a 'fortress-like' barrier that impairs drug delivery to tumour cells [179]. However, there is now an increasing body of evidence that suggests that, in addition to inanimate structural effects, the desmoplastic reaction is in fact a dynamic contributor to pancreatic cancer progression through active modulation of key signalling events regulating both tumour cell proliferation and invasion. As discussed above and summarized in Figure 4, the intricate interplay between collagen, MT1-MMP and TGF- $\beta$  promotes pancreatic cancer invasion, represses tumour-suppressive microRNAs, and provides critical survival and repair signals to mediate

chemoresistance in PDAC cells. Feed-forward amplification loops involving collagen, MT1-MMP and TGF- $\beta$  signalling further promote the establishment and maintenance of the desmoplastic reaction. On the basis of results from several *in vivo* studies, targeting the key collagenase MT1-MMP may help to not only limit pancreatic cancer progression, but also paradoxically attenuate the pronounced stromal reaction that is both a characteristic descriptive hallmark of the disease and now an increasingly recognized functional component of pancreatic cancer's aggressive molecular and clinical trajectory. A highly selective anti-MT1-MMP antibody has been shown to inhibit growth and metastasis in a mouse model of breast cancer [180]. Evaluating this antibody in similar mouse models of pancreatic cancer will provide proof-of-principle evidence that targeting MT1-MMP can prevent pancreatic cancer growth and invasion. Moving forward, a key objective in the effective treatment of pancreatic carcinoma is thus the development of strategies to breach the stromal fortress that heretofore has not only enabled metastatic forays into normal tissue, but also acted as a nearly impenetrable barrier for therapeutic drug delivery.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Abbreviations used:

<b>AP-1</b>	activator protein 1
<b>CDK</b>	cyclin-dependent kinase
<b>ECM</b>	extracellular matrix
<b>EMT</b>	epithelial–mesenchymal transition
<b>ERK</b>	extracellular-signal-regulated kinase
<b>HCC</b>	hepatocellular carcinoma
<b>HMGA2</b>	high-mobility group A2
<b>ILK</b>	integrin-linked kinase
<b>LAP</b>	latency-associated peptide
<b>MDCK</b>	Madin–Darby canine kidney
<b>MMP</b>	matrix metalloproteinase
<b>MT1-MMP</b>	membrane-type 1 MMP
<b>PDAC</b>	pancreatic ductal adenocarcinoma

<b>PDGF-<math>\beta</math></b>	platelet-derived growth factor- $\beta$
<b>PI3K</b>	phosphoinositide 3-kinase
<b>PTEN</b>	phosphatase and tensin homologue deleted on chromosome 10
<b>R-Smad</b>	receptor-associated Smad
<b>siRNA</b>	small interfering RNA
<b>TGF-<math>\beta</math></b>	transforming growth factor- $\beta$
<b>T<math>\beta</math>R</b>	TGF- $\beta$ receptor
<b>TIMP</b>	tissue inhibitor of metalloproteinases
<b>VSMC</b>	vascular smooth muscle cell

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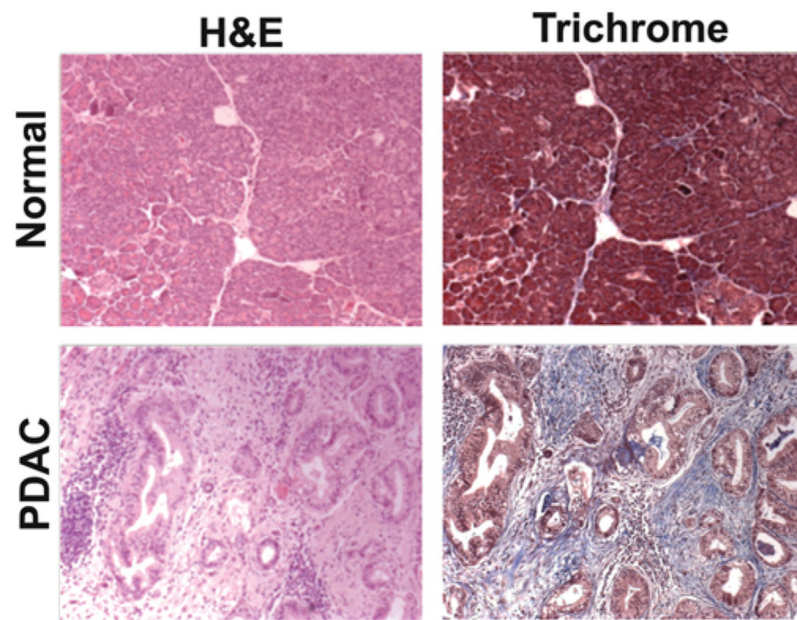
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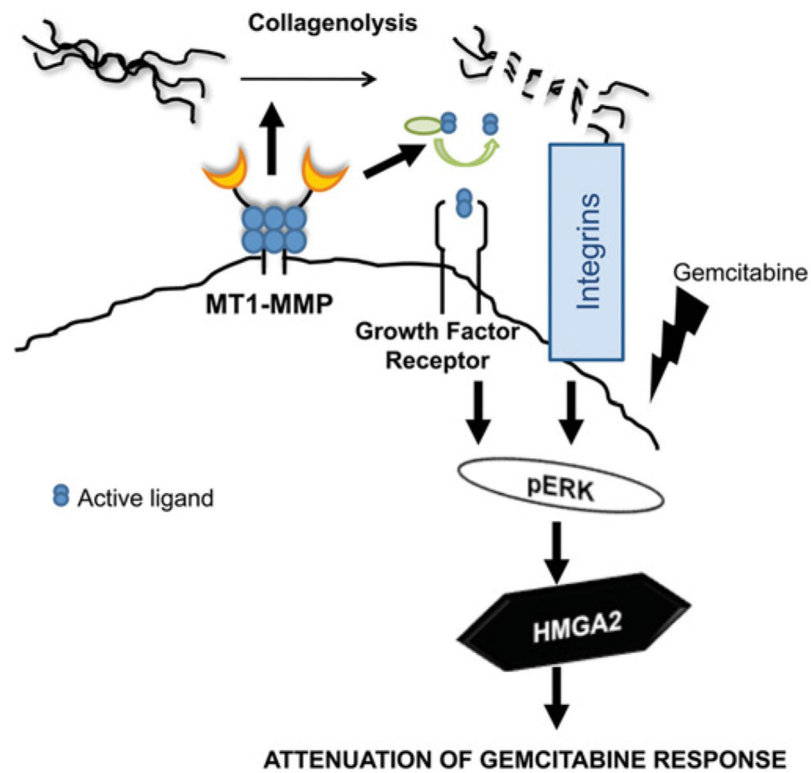
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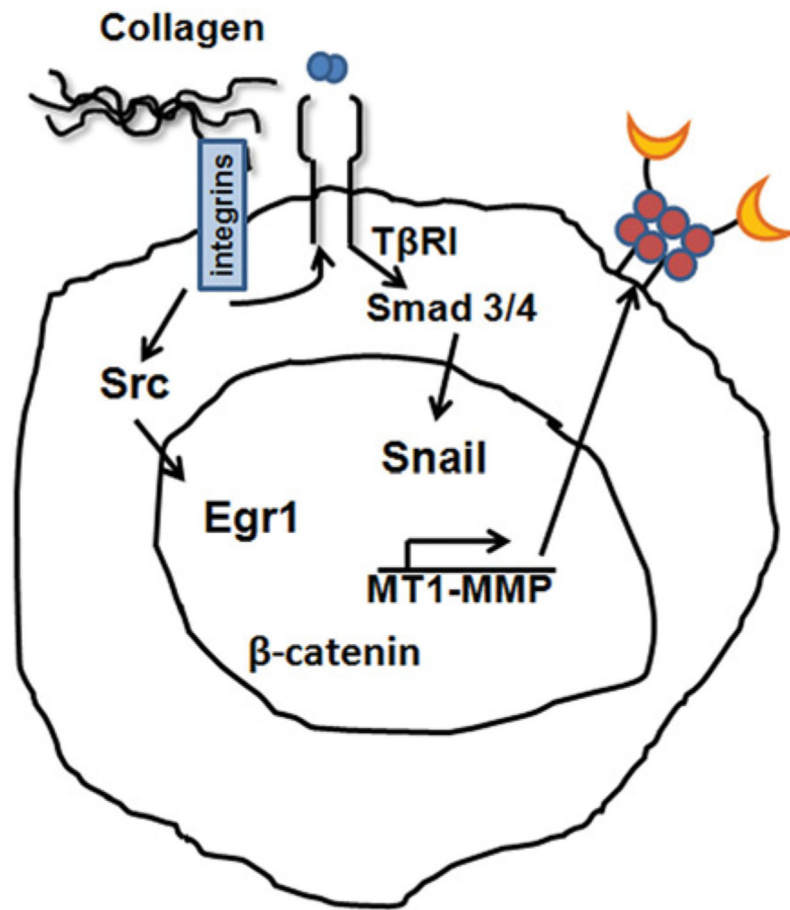
**Figure 1. Desmoplastic reaction and pancreatic cancer**

H&E (haematoxylin and eosin) and trichrome stains of normal pancreas and a human PDAC tumour. Note the increased fibrosis (blue staining) in the PDAC tumour compared with the normal pancreas.



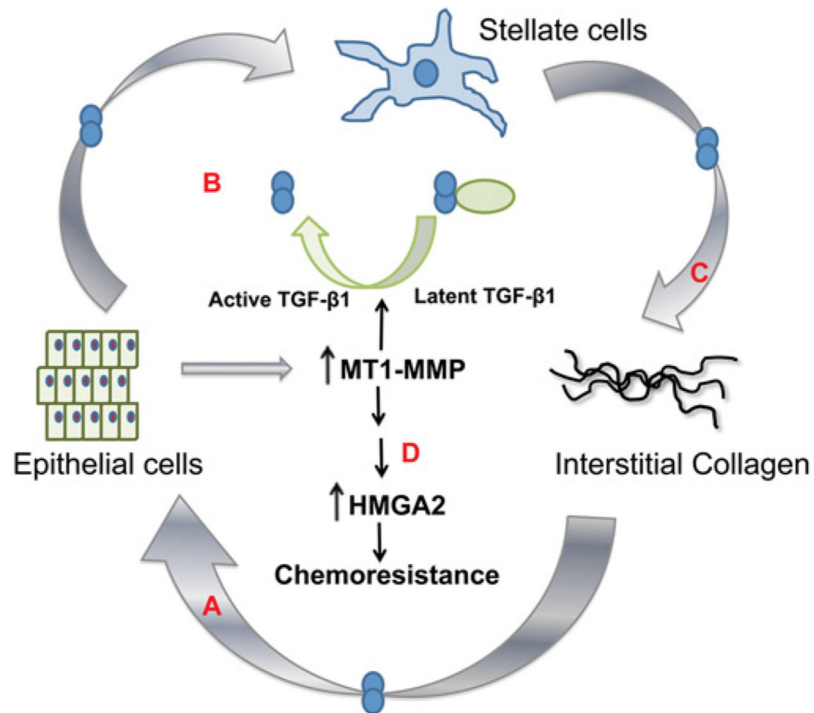
**Figure 2. MT1-MMP in three-dimensional collagen contributes to chemotherapy resistance by increasing ERK1/2 activity and HMG2 levels**  
 MT1-MMP through proteolytic processing of collagen or activation of growth factors modulates integrin and/or growth factor receptor signalling to enhance ERK1/2 phosphorylation and HMG2 expression in pancreatic cancer cells. This signalling pathway functions to attenuate the effect of gemcitabine chemotherapy in the three-dimensional collagen microenvironment.





**Figure 3. Collagen regulation of MT1-MMP mRNA expression**

Pancreatic cancer cells in three-dimensional collagen up-regulate MT1-MMP through activation of the TGF- $\beta$ /T $\beta$ RI/Smad3/Snail pathway. MT1-MMP can also be induced in the collagen microenvironment through the  $\beta$ 1-integrin/Src/Egr1 signalling pathway. Although  $\beta$ -catenin signalling can regulate MT1-MMP, it is not known whether collagen regulation of MT1-MMP also involves  $\beta$ -catenin signalling.



**Figure 4. A feed-forward amplification loop involving collagen, MT1-MMP and TGF- $\beta$**   
**(A)** Pancreatic cancer cells in the collagen microenvironment up-regulate TGF- $\beta$  signalling to induce MT1-MMP expression. **(B)** Expression of MT1-MMP in cancer cells can in turn cause stellate cell activation also through increased TGF- $\beta$  signalling. **(C)** The activated stellate cells deposit more collagen, resulting in a further increase in MT1-MMP expression in the cancer cells. As detailed in the text, this forward amplification loop contributes to pancreatic cancer invasion and generation of the desmoplastic reaction. **(D)** MT1-MMP expression in the collagen microenvironment also contributes to chemoresistance, in part, through up-regulation of HMGA2 (see also Figure 2). An animated version of this Figure is available at <http://www.BiochemJ.org/bj/441/0541/bj4410541add.htm>.

**Table 1**

Examples of microRNA modulating fibrosis or chemoresistance

MicroRNA	Organ/cancer	Phenotype	Target (direct or indirect)	Reference(s)
<i>let-7d</i>	Lung	Decreases fibrosis	Inhibits <i>COL1A1</i>	[166]
<i>let-7g</i>	HCC	Decreases fibrosis; reduces migration and cell growth	Inhibits <i>COL1A2</i>	[167]
<i>miR-29b</i>	Liver	Reduces collagen production	Inhibits <i>COL1A1</i> , <i>COL4A5</i> and <i>COL5A3</i>	[168,169]
<i>miR-200</i>	Kidney	Reduces fibrosis	Inhibits collagen I and III	[172]
<i>miR-192</i>	Kidney	Promotes fibrosis and collagen I downstream of TGF- $\beta$ /Smad3	Induces collagen I	[173]
<i>miR-21</i>	PDAC	Gemcitabine resistance	Inhibits PTEN and Bax; induces Bcl-2	[175,176]
<i>miR-21</i>	Bladder cancer	Doxorubicin resistance	Inhibits PTEN	[177]