Inflammation and EMT: an alliance towards organ fibrosis and cancer progression

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Introduction—EMT in cancer and fibrosis

Although inflammation has been associated with the progression of chronic kidney disease and cancer for decades, the molecular mechanisms involved have remained elusive until recently (Cordon-Cardo & Prives, 1999; Kalluri & Neilson, 2003; Mantovani et al, 2008). The epithelial to mesenchymal transition (EMT) now takes centre stage as the convergence point between inflammation and the progression of degenerative fibrotic diseases and cancer. This cellular process is characterized by the loss in cell polarity and a change in cell shape from cuboidal to fibroblastoid, the downregulation of epithelial markers and the upregulation of mesenchymal markers. The cell acquires the capacity to degrade the basement membrane and migrate through the extracellular matrix to populate different territories either during embryonic development or cancer progression, or to adopt a profibrotic myofibroblast nature in the renal interstitial space (Acloque et al, 2009; Kalluri & Neilson, 2003; Kalluri & Weinberg, 2009).

Betty Hay was the first to coin the term ‘epithelial to mesenchymal transformation’ in embryos (Hay, 1968), as well as later describing this cellular behaviour during migration (Hay, 1990) and the importance of the transient nature of this process (Hay, 1991; Fig 1A). Ironically, important elements in the control of the inflammatory response and in the induction of tumour cell death such as TGF-β1 and hypoxia can also act as potent inducers of EMT in an inflammatory microenvironment. EMT events have been very well defined during embryonic development for many years (reviewed in Acloque et al, 2009). However, the significance of the EMT in cancer progression has been debated until recently. The main discussion regarding the role of EMT in cancer came from the difficulty in finding morphological evidence from clinical samples, due to the lack of specific markers for many years and the sometimes focal nature of the event. Nevertheless, more than 100 years ago, both Santiago Ramón y Cajal and Hugo Apolant noticed what we would definitely consider an EMT in both human and mouse breast cancer (Fig 1B,C). Indeed, the description of the breast tumours by Cajal in his ‘Manual of Anatomopathology’ is premonitory of the implication of the EMT as the first step in the metastatic cascade. He referred to undifferentiated breast carcinomas as follows: ‘The epithelial islands are not surrounded by a basement membrane... We shall mention the fusiform, pear-like and star-like forms... The cells are not attached to each other... This explains their invasive ability, since free of intercellular cement, they can migrate through the connective tissue’ (Ramon y Cajal, 1890). When we read
‘cement’ in Cajal’s words we immediately think of E-cadherin, the molecule that maintains epithelial cells together and that we now know is the main target repressed by the EMT inducers. Figure 1B and C shows fragments of Cajal’s and Apolant’s drawings illustrating human and mouse breast carcinomas. The reversibility of the EMT process has been well described in the context of embryonic development, where mesenchymal to epithelial processes (MET) are fundamental for the differentiation of tissues and organs once the embryonic migratory cells have reached their destination. Indeed, the MET events certainly take place during metastasis formation as suggested by cell morphology and the re-expression of E-cadherin expression (Bukholm et al, 2000). This EMT–MET double switch for metastasis formation has been discussed by Brabletz et al (2001) while providing a convincing illustration of an EMT process at the invasive front of human colon carcinoma (Fig 2A).

Fibrosis is characterized by the presence of an excess of fibrous connective tissue in an organ, and in particular by an excessive deposition of collagen I. It can be the result of a reparative or reactive process that can occur in the lung, the liver, the heart and the kidney among other organs. Under chronic pathological situations, fibrosis progresses to advanced states that lead to defective organ function and final organ failure. In this review we will focus on renal fibrosis as it has been very well studied at the cellular level and the mechanisms leading to the development of fibrosis are thought to be very similar in different organs. Renal fibrosis is the link between the progressive loss of renal function and primary diseases such as glomerulonephritis, diabetes, toxic injury, congenital abnormalities, urinary tract obstruction and chronic rejection of transplanted kidneys (Kalluri & Neilson, 2003; Liu, 2006; Vongwiwatana et al, 2005). Renal fibrosis has been associated with the activation of interstitial fibroblasts to give rise to collagen secreting myofibroblasts. However, different studies have shown that in addition, myofibroblasts can also originate from renal tubular epithelial and endothelial cells that undergo EMT in mouse models of renal fibrosis (Iwano et al, 2002; Zeisberg et al, 2008). Furthermore, the aberrant activation of Snail1, a well-known EMT inducer leads to the appearance of renal fibrosis and renal failure in transgenic mice (Boutet et al, 2006). Interestingly, high Snail1 expression and evidence of EMT has also been found in the kidneys of patients with renal fibrosis (Boutet et al, 2006; Jinde et al, 2001; Rastaldi et al, 2002).

In this review, we will discuss the dual role of TGF-β1 and hypoxia in fibrotic tissues and in the tumour microenvironment. We propose that in the context of a chronic inflammatory
condition, TGF-\(\beta\)1 and hypoxia reactivate EMT developmental programmes that converge in the activation of NF-κB also induced by the inflammatory cytokines and oxidative stress. These EMT programmes, in an unsuccessful attempt to repair the injured tissue, turn to a sinister role and lead to the destruction of epithelial homeostasis and accumulation of extracellular matrix in fibrosis, and to the progression of carcinomas towards the metastatic state.

**Inflammation as an inducer of fibrosis and cancer**

**Inflammation and cancer**

The link between cancer and inflammation has been recognized for decades, as strong associations have been established between chronic inflammatory conditions and tumourigenesis (Cordon-Cardo & Prives, 1999). As such, ulcerative colitis, chronic gastritis, hepatitis and chronic pancreatitis, and their respective relationships with colon, gastric, liver and pancreatic carcinomas exemplify the close connection between inflammation and tumour appearance. Furthermore, non-steroidal anti-inflammatory drugs reduce both the risk of cancer development and its mortality (Mantovani et al, 2008).

**Renal fibrosis as an inflammatory disease**

Renal fibrosis is a paradigmatic example of organ fibrosis towards degenerative organ disease and final stage kidney damage leading to the destruction of the tissue and death from renal failure. Obstructive nephropathy (ON) in humans, either due to congenital or acquired obstruction of the urinary tract, is the first primary cause of chronic renal failure (CRF) in children and a major cause of kidney failure in adults (Klahr & Morrissey, 2002; Smith et al, 2007). In ON, the evolution of renal disease is similar to that occurring in polycystic kidney disease or renal transplant rejection. Moreover, in its final fibrotic phase, ON is also very similar to the fibrosis secondary to glomerulonephritis, diabetes or hypertension (Chevalier, 2006; Chevalier et al, 2009). Indeed, long-term ON is characterized by tubular atrophy and the appearance of fibrosis (Fig 2B). Studies of ON have benefited from the existence of a recognized experimental model, the unilateral ureteral obstruction (UUO), in mice. UUO does not compromise the life of the animal, as the contralateral kidney maintains or even augments its function offering functional compensation (Klahr & Morrissey, 2002). Since UUO reproduces all the hallmarks of ON and it generates progressive renal fibrosis, it has become the standard experimental model to study the causes and mechanisms underlying tubulointerstitial fibrosis (Chevalier et al, 2009). Even though ON is not an immune disease, it has a major inflammatory component, characterized by the overexpression of inflammatory genes, the release of pro-inflammatory cytokines, the activation of NF-κB and the infiltration of macrophages and lymphocytes (Diamond, 1995; Esteban et al, 2004; Misseri et al, 2005; Silverstein et al, 2003). Similarly, fibrosis can also be considered as the end result of chronic inflammatory reactions induced by a variety of stimuli including persistent infections, autoimmune reactions, allergic responses, chemical insults,
radiation, tissue injury and normal ageing (Kalluri & Neilson, 2003). Therefore, ON and UUO can be used as a reference for the development of fibrosis, in part because inflammatory responses have been analysed in depth in this model.

NF-κB activation in fibrosis and cancer

After renal obstruction, an increase in the expression of the prototypical pro-inflammatory cytokines tumour necrosis factor (TNF-α) and interleukin-1 (IL-1) is crucial for the induction of NF-κB, the major inflammatory response pathway, and for recruitment of inflammatory cells to the obstructed kidney (Meldrum et al, 2006; Metcalfe et al, 2006; Misseri et al, 2005; Silverstein et al, 2006; Yamagishi et al, 2001). An increase in leukocyte infiltration, especially macrophages and T lymphocytes, is detected as early as 4–12 h after ureteral obstruction, and it continues to increase over the following days (Diamond, 1995). These cells play a central role in the renal inflammatory response to UUO as the progression of renal injury is closely associated with their accumulation and blocking their recruitment protects against interstitial fibrosis (Lange-Sperandio et al, 2007). The recruitment of circulatory leukocytes is mediated by several mechanisms including the expression of pro-inflammatory, profibrotic and chemoattractant cytokines in the kidney and the corresponding leukocyte chemotactic response to those chemokines (Ferenbach et al, 2007).

Once activated, NF-κB generates a loop that maintains the inflammatory signals as it controls the expression of genes encoding pro-inflammatory cytokines (e.g. IL-1, IL2, IL-6, TNF-α, etc.), chemokines (e.g. IL-8, MIP-1 α, MCP-1, RANTES, eotaxin, etc.), adhesion molecules (e.g. ICAM, VCAM, E-selectin), inducible enzymes (COX-2 and iNOS), growth factors, some of the acute phase proteins and immune receptors, all of which play critical roles in controlling most inflammatory processes (Blackwell & Christman, 1997; Fig 3). As inflammation plays an important role in fibrosis, inhibition of NF-κB has a big impact on the development of chronic kidney disease. Indeed, NF-κB inactivation prevents inflammatory injury and diminishes the expression of inflammatory genes after UUO (Esteban et al, 2004; Miyajima et al, 2003). Similarly, increasing the levels of the endogenous inhibitor of NF-κB l-κB, or inhibiting the pathway by the administration of curcumin, reduces renal fibrosis and macrophage influx following UUO or protects against interstitial inflammation in ON, respectively (Kuwabara et al, 2006; Tashiro et al, 2003).

Like the injured kidney, inflammation or infection in the context of a neoplasia can trigger the activation of NF-κB, the main transducer of inflammatory responses, which in turn induces the secretion of a plethora of inflammatory cytokines (Fig 3). These cytokines recruit inflammatory cells and, at the tumour–stroma interface, tumour cells together with tumour-associated macrophages (TAMS) and cancer-associated fibroblasts (CAFs) again converge in producing NF-κB and the hypoxia-inducible factor (HIF1-α), generating a microenvironment capable of driving tumour progression (Mantovani et al, 2008). HIF-1 is a potent inducer of EMT in the kidney and also in cancer cells in general (see Box 1 and Fig 3) and the inflammation-mediated increase in NF-κB expression has been associated with several aspects of cancer development, mainly resistance to apoptosis and increased angiogenesis, with some connections to invasiveness (Karim, 2006).

NF-κB can also directly activate the expression of potent EMT inducers, including Snail1 and Zeb factors (Bachelder et al, 2005; Barbera et al, 2004; Chua et al, 2007; Julien et al, 2007; Kim et al, 2007) (Fig 3). Thus, there seems to be an important connection between cancer, inflammation and EMT. This connection has been reinforced recently by the finding that TNF-α induces Snail1 promoter activity and EMT in cancer cells (Dong et al, 2007) and that it can also stabilize Snail1 protein (Wu et al, 2009). Snail1 is a highly unstable protein, targeted for degradation by phosphorylation and ubiquitylation mediated by GSK-3β and SCFβ-Trcp, respectively (Zhou et al, 2004). TNF-α induces Snail1 protein stabilization by activating the expression of the COP9 signalosome 2 (CNS2), which in turn inhibits the binding of Snail1 to GSK-3β and β-Trcp (Wu et al, 2009; Fig 3).
Interestingly, both transcriptional and post-transcriptional mechanisms act through the induction of the main target of TNF-α in the inflammatory response, NF-κB.

The link between TNF-α and Snail factors points to the EMT as the molecular and cellular explanation for the association between inflammation, fibrosis and tumour progression. As mentioned above, EMT events are now directly associated with invasion and metastasis in cancer progression. Elegant in vivo imaging studies have shown that carcinoma cells migrate from mouse primary tumours through a process of EMT and that this process is dependent on an inflammatory microenvironment provided by the TAMs and other stromal cells such as the CAFs (Condeelis & Segall, 2003) (Fig 3). TAMs and stromal cells in general are essential for angiogenesis, extracellular matrix remodelling and the inflammatory response associated with cancer (Condeelis & Pollard, 2006). Stromal cells secrete TNF-α which, as in renal cells in the obstructed kidney, induces NF-κB in the tumour cells. The significance of NF-κB activation in cancer cells is evident from the results obtained in inflammation-associated models of cancer (Karin & Greten, 2005; Pikarsky et al., 2004). Selective inactivation of NF-κB in these tumours led to deficient tumour development. Importantly, at least in the liver, activation of NF-κB was not important in the early stages of tumour development, but was crucial for malignant conversion and for the survival of tumour cells. This combined effect is very reminiscent of that produced by the EMT inducers and Snail in particular as, in addition to triggering EMT, Snail-expressing cells are resistant to the cell death induced by the removal of survival factors, by genotoxic stress, chemotherapy and immunotherapy and even by TNF-α (reviewed in Barrallo-Gimeno & Nieto, 2005; see also Kudo-Saito et al., 2009; Vega et al., 2004). In addition to the activation of Snail1 by NF-κB, it should be noted that angiotensin...
In summary, while some steps of the pathways involved may be tissue or disease specific, it is clear that in both cancer and fibrosis the concerted action of several cytokines as IL-1β and TNF-α plus Ang II together with oxidative stress and hypoxia converge on the activation of NF-κB, a major inducer of the inflammatory response.

**Yin and Yang of TGF-β1**

In addition to the secretion of inflammatory cytokines, many organs and tissues release TGF-β1 after an injury, as seen in the lung, liver, heart, kidney and bone (Devescovi et al., 2008; Frangogiannis, 2008; Kaneto et al., 1993; Kisseleva & Brenner, 2008; Gharave-Kermani et al., 2009; Gressner et al., 2008). TGF-β1 is considered a major anti-inflammatory cytokine, as TGF-β1 knockout mice suffer from a lethal multifocal inflammatory disease (Kulkarni et al., 1993; Letterio & Roberts, 1998) and the blockage of TGF-β1 signalling in T cells or in bone marrow results in similar multifocal inflammatory responses (Gorelik & Flavell, 2000; Shull et al., 1992). However, TGF-β1 has been known for many years to display a schizophrenic behaviour in cancer. While it can act as a tumour suppressor at early tumour stages, it can later contribute to the malignant progression by promoting invasion and metastasis (Massague, 2008). In the kidney too, although TGF-β1 acts as an anti-inflammatory cytokine to heal the injured kidney, it also has negative effects on the development of chronic renal disease as it is an important fibrogenic agent. Indeed, the role of TGF-β1 as a promoter of tumour progression and profibrotic agent is associated with its ability to act as a potent activator of EMT inducers (Acloque et al., 2009). The Yin and Yang of TGF-β1 in both cases are explored below.
TGF-β1 in cancer: tumour suppressor and promoter of invasion and metastasis

There are paradigmatic examples illustrating how TGF-β1 can impair tumourigenesis while promoting metastasis (Siegel et al, 2003, reviewed in Massague, 2008). In normal cells, TGF-β1 can maintain tissue homeostasis by suppressing cell proliferation and by inducing apoptosis. Tumour cells evade this response by losing the ability to respond to the cytostatic and apoptotic effects through different mechanisms (Shi & Massague, 2003). In addition, TGF-β1 helps cancer cells to acquire the ability to invade and disseminate by inducing the EMT process. Thus, TGF-β1 is a potent EMT inducer in tumours albeit it is also initially secreted to control pathological proliferation and inflammation.

The study of the molecular pathways that drive the differential response to TGF-β1 continues to be a very active field of research and recent findings indicate that both genetic and epigenetic mechanisms are involved. On one hand, the methylation state of the platelet-derived growth factor (PDGF) gene limits the proliferative or cytostatic effect of TGF-β1 in gliomas (Bruna et al, 2007) and the concerted action of TGF-β1 with oncogenic Ras and mutant p53 sequester p63 in a complex that inactivates its tumour suppressor effects (Adorno et al, 2009). This indicates that two molecules frequently mutated in cancer, p53 and Ras, can divert the response to TGF-β1 towards the promotion of metastasis. Indeed, TGF-β1 and oncogenic Ras cooperate in the induction of Snail1 expression and EMT in epithelial cells (Peinado et al, 2003). Interestingly, NF-κB is required for the induction and maintenance of the EMT by TGF-β1 in mammary epithelial cells overexpressing the Ras oncogene (Huber et al, 2004), explaining the role of NF-κB as a modulator of TGF-β1-induced EMT in addition to its role as a direct activator of the EMT inducers.

But what are the sources of TGF-β1 in the tumours? Again, as in the injured organs, TGF-β1 is produced by cancer cells and, particularly, by stromal cells in an attempt to control inflammation. TNF-α can induce TGF-β1 in lung fibroblasts (Sullivan et al, 2009) and TGF-β1 can also activate the resident fibroblasts to convert them into myofibroblasts, called cancer-associated fibroblasts in the contexts of tumours (CAF) and associated with tumour invasion (Mickey & Ostman, 2005). Some of these ‘myofibroblastic’ CAFs are surely the result of the EMT undergone by carcinoma cells, as EMT is now recognized as an important event in carcinoma progression.

TGF-β1 in fibrosis: anti-inflammatory and healing factor versus profibrotic factor

The release of TGF-β1 by renal or by infiltrating cells in the damaged kidney moderates the inflammatory reaction and helps to heal the damaged tissue. Among its anti-inflammatory functions, TGF-β1 antagonizes the pro-inflammatory cytokines IL-1 and TNF-α in glomerular disease, and it is a prominent macrophage deactivator during kidney injury (Kitamura & Suto, 1997). Renal TGF-β1 mRNA expression increases considerably after the onset of obstruction (Kaneto et al, 1993), and its concentration increases in the plasma of patients with obstruction due to ureteral calculi and renal fibrosis (Liu, 2006; Vuruskan et al, 2005). TGF-β1 expression is induced by Ang II (Wolf, 2006), upon increased stretch of tubular epithelial cells (Quinlan et al, 2008), and it is also produced by interstitial fibroblasts and infiltrating macrophages (Diamond et al, 1998; Ding et al, 1993). As already mentioned, in addition to its anti-inflammatory effects, TGF-β1 also acts as a profibrotic agent. The profibrotic effect of TGF-β1 is achieved by increasing the synthesis of matrix proteins and the expression of proteinase inhibitors, including plasminogen activator inhibitor-1 (PAI-1), and by decreasing that of ECM-degrading proteins such as collagenase (Chevalier, 2006). Therefore, sustained aberrant expression of TGF-β1 results in the pathological accumulation of extracellular matrix material in both the glomerulus and interstitial compartments (Bottinger, 2007). TGF-β1 also has some pro-inflammatory properties, as it functions as a chemo-attractant for leukocytes (Wahl et al, 1987) and induces COX-2 in mesangial cells (Rodrı´guez-Barbero et al, 2006). This profibrotic activity of TGF-β1 reflects its behaviour as a potent inducer of EMT, which contributes to the conversion of renal epithelial cells into myofibroblasts in the context of the injured kidney. In addition, part of the TGF-β1-induced EMT programme promotes the remodelling of the cell contacts with the basal membrane by activating the matrix metalloproteases MMP2 and MMP9 (Li et al, 2003; Strutz et al, 2002), leading to the degradation of the collagen type IV component of basement membrane. Thus, the molecule secreted to control inflammation has an alter ego and it promotes the development of fibrosis.

TGF-β1, myofibroblasts and EMT

Myofibroblasts play a major role in interstitial fibrosis as they are the main source of matrix proteins. They originate from several sources, including bone marrow cells, resident interstitial fibroblasts, vascular pericytes, as well as endothelial and tubular epithelial cells (reviewed in Grande & Lopez-Novoa, 2009). As mentioned above, TGF-β1 contributes to the formation of myofibroblasts through the activation of resident fibroblasts and by inducing the transition to mesenchyme of epithelial and endothelial cells via EMT or EndMT, respectively (Zavadil & Böttinger, 2005; Zeisberg & Kalluri, 2008; Zeisberg et al, 2008). Indeed, the transformation of tubular and endothelial cells to mesenchymal cells contributes more than 60% of the myofibroblast population in the obstructed kidney after UUO (Iwano et al, 2002; Zeisberg et al, 2008), and this process plays a major role in tubulo-interstitial fibrosis (Kalluri & Neilson 2003). The induction of EMT by TGF-β1 also explains the fibrotic effect of Ang II, as this anti-inflammatory factor also induces the synthesis of TGF-β1 and its receptors in tubular epithelial cells (Wolf, 2006). Accordingly, the infusion of Ang II into mice subjected to UUO significantly increases their collagen content (Ma et al, 2003). The induction of EMT by TGF-β1 in the renal cells takes several days and requires the interplay of many pathways, involving SMAD proteins (Li et al, 2002), the integrin-linked kinase (Li et al, 2003) and Rho-family GTPases (Bhowmick et al, 2001; Masszi et al, 2003). β-Catenin is another key element in the myofibroblast transition as it synergizes with TGF-β1 in
activating the α-smooth muscle actin (αSMA) gene and protein expression (Masszi et al., 2004). The processes triggered by TGF-β1 converge in the activation of the so-called EMT inducers, which are transcription factors capable of eliciting this dramatic phenotypic change that converts epithelial cells into activated ‘myofibroblast-like’ cells. As in cancer, among the transcription factors involved in the induction of EMT, Snail factors have been associated with renal fibrosis and significantly, TGF-β1 is the most potent inducer of Snail transcription (Fig 3). The expression of these genes increases in the obstructed kidney after UUO (Sato et al., 2003; Tan et al., 2006; Yoshino et al., 2007) and Snail1 activation is sufficient to induce EMT and all the hallmarks of kidney fibrosis in adult mice (Boutet et al., 2006). Snail1 directly represses E-cadherin transcription and that of the renal-specific Cadherin16 gene through the direct repression of HNF-1β gene expression. Snail also activates, albeit indirectly, the transcription of mesenchymal genes, such as vimetin and αSMA, and it promotes collagen I synthesis and deposition (Boutet et al., 2006). Interestingly, Snail factors are strongly upregulated in fibrotic kidneys from patients subjected to nephrectomy due to urinary obstruction and kidney failure (Boutet et al., 2006). Altogether, data from many laboratories indicate that Snail factors are crucial mediators of TGF-β1-induced EMT.

Concluding remarks and perspectives

From all the work carried out in the context of organ fibrosis and cancer we can conclude that the pathways and players identified that lead to pathological EMTs are basically the same. Both in cancer and fibrosis TGF-β1, TNF-α and hypoxia cooperate in the triggering of EMT, converging in the induction of Snail activity through different mechanisms among which the activation of NF-κB seems to play a central role. This can be regarded as the reactivation of a developmental programme designed to endow cells with migratory and invasive properties and with amazing survival properties, crucial for migratory cells to reach their sometimes far destinations within the embryo (Acloque et al., 2009). This developmental programme, firstly reactivated to control the inflammatory response and to heal the injured tissue, is corrupted in the context of chronic inflammation and cancer, favouring the survival of injured or malignant cells and stimulating the production of pro-inflammatory cytokines in the tumour mass or in the organ interstitial space. The cooperation between TGF-β1 and the pro-inflammatory cytokines generates a microenvironment that favours the generation of autoregulatory loops to reinforce the EMT programme (Fig 3). Not only Ang II, TNF-α, ROS and hypoxia converge in the induction of Snail, but also Ang II and TNF-α can induce TGF-β1 at the transcriptional level (Sullivan et al., 2009), the most potent Snail inducer. In addition, Snail1 itself also seems to upregulate the expression of pro-inflammatory mediators such as several ILS (IL-1, IL-6, IL-8; Lyons et al., 2008). It will be interesting to determine whether other EMT inducers are also activated in the context of this inflammatory microenvironment, which seems to be very likely, given that both TGF-β1 and hypoxia can induce the expression of Twist and Zeb transcription factors.

The relationship between inflammation and Snail activity has been established in cancer cells from different origins, indicating that this is a general mechanism in carcinoma progression. While the majority of studies on fibrosis have been carried out in the kidney, the same pathways are likely to operate during other fibrotic processes, including those in the liver and the lung, especially since there is evidence of EMT in hepatic fibrosis (Kim et al., 2006; Zeisberg et al., 2007a). A similar process also occurs when endothelial cells undergo an EMT during cardiac or renal fibrosis (EndMT; Zeisberg et al., 2007b, 2008) as well as during mesothelial fibrosis in patients subjected to peritoneal dialysis where the NF-κB/Snail1 axis has already been described (Strippoli et al., 2008).

In summary, the relationship between inflammation and EMT seems to be a conserved feature in the progression of organ fibrotic diseases and cancer, suggesting that specific anti-EMT drugs should be considered to treat these conditions in combination with anti-inflammatory therapies. With respect to therapies targeting the EMT programme, small molecule inhibitors or antibodies directed against TGFβR have been effective in preclinical and clinical trials, and neutralizing antibodies against TGF-β1 are in Phase 1 clinical trials for renal cell carcinoma and pulmonary fibrosis (Chua et al., 2008). Treatment with BMP7, an inhibitor of TGF-β1 signalling, is a promising strategy to treat fibrosis as it can reverse TGF-β1-induced renal fibrosis in mice (Zeisberg et al., 2003). It should be noted that since signalling molecules such as TGF-β1 or hypoxia trigger complex transduction cascades with deleterious but also beneficial outcomes, inhibiting the EMT inducers rather than the extracellular signalling molecules or their receptors could provide a more specific way to fight both fibrosis and cancer progression. However, EMT inducers are transcription factors, and thus very difficult to target. While RNA interference approaches are promising specific reagents, these are still early days for their wide use due to their low stability and, more importantly, due to the current low efficiency in their cell targeting and intracellular delivery. Finally, as the same signalling pathways can act as promoters or suppressors of fibrosis and cancer, better understanding of the cellular response that depends on the context and stage of the disease is crucial to design more advanced therapies.

Pending issues

- The development of additional in vivo models to assess the full significance of EMT in cancer and organ fibrosis.
- The characterization of the full complement of EMT inducers activated in organ fibrosis.
- The development of optimized antibodies specific for the EMT inducers that can work on histopatological sections.
- The development of therapies directed towards the inhibition of the EMT process.
- The optimization of methods for the delivery of RNA interference reagents.
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