Exploiting inflammation for therapeutic gain in pancreatic cancer

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Pancreatic ductal adenocarcinoma (PDAC) is an aggressive malignancy associated with <5% 5-year survival, in which standard chemotherapeutics have limited benefit. The disease is associated with significant intra- and peritumoral inflammation and failure of protective immunosurveillance. Indeed, inflammatory signals are implicated in both tumour initiation and tumour progression. The major pathways regulating PDAC-associated inflammation are now being explored. Activation of leukocytes, and upregulation of cytokine and chemokine signalling pathways, both have been shown to modulate PDAC progression. Therefore, targeting inflammatory pathways may be of benefit as part of a multi-target approach to PDAC therapy. This review explores the pathways known to modulate inflammation at different stages of tumour development, drawing conclusions on their potential as therapeutic targets in PDAC.

Pancreatic ductal adenocarcinoma (PDAC) is one of the most lethal and aggressive of all malignancies, and as the 5th most common cause of cancer death in the United Kingdom represents a significant therapeutic challenge. Patients with potentially resectable disease are in the minority (10–15%) because of extensive local spread or metastatic disease at presentation (Sener et al, 1999). Despite surgery, the 5-year survival rate is a dismal 3–5% due to secondary recurrence.

The European Study of Pancreatic and Ampullary Cancer (ESPAC) trials have consistently demonstrated a modest survival benefit associated with post-operative adjuvant therapy, with fluorouracil and gemcitabine equally effective (Neoptolemos et al, 2010). Unfortunately, although a subset of patients who receive gemcitabine have tumours responsive to these drugs, a significant number develop resistance (Liss and Thayer, 2012). Despite similar presentation and histological appearance, these tumours are biologically diverse and exhibit complex molecular and cellular heterogeneity, however, aggressive progression is common to all (Vincent et al, 2011). Indeed, recent elegant lineage tracing studies in mice have identified that cells may disseminate from pancreatic lesions even before the primary tumour is histologically evident, leading to a call for new approaches to PDAC therapy (Rhim et al, 2012; Tuveson and Neoptolemos, 2012). Management algorithms for PDAC must change, taking into account tumour biology, and stratifying therapy on an individual patient basis while pursuing strategies to combat recurrence in patients following surgical resection. The dependence of PDAC on certain crucial inflammatory mediators suggests that targeting of these central roles may hold hope for effective therapies.

The following review focuses on the failure of tumour immunosurveillance mechanisms in PDAC initiation and the accumulation in the pancreas of immunosuppressive cells that permits tumour progression, and discusses how failure of immunosuppression may be addressed therapeutically. We describe how tissue damage combined with oncogene activation (KRAS), loss of tumour suppressor genes (TSGs) (TP53 and SMAD4), deregulation of chemokines (CXCR2 ligands) and cytokines (interleukin (IL)-6, IL-1α), activation of downstream effectors (STAT3, NF-κB), and concurrent leukocyte recruitment by the tumour are key events in both initiation and progression of PDAC. Throughout, we consider the potential of these inflammatory mediators as therapeutic targets in PDAC taking into account the results of preclinical and clinical research to date.

**IMMUNE INFILTRATION DURING PDAC INITIATION AND PROGRESSION**

In healthy individuals, the adaptive immune system is responsible for tumour immunosurveillance, through which the host can identify, mount a response to, and destroy tumour cells.
Tumorigenesis represents a failure of adaptive immunity, as in order for malignant transformation to progress unchecked tumour cells must escape this surveillance. The immune response to cancer is a dynamic process and can also initiate pathways that are in fact pro-tumorigenic. Even then a fine balance exists, as tumour progression is complex, reliant on interactions between the tumour and its activated microenvironment (Baumgart et al, 2011).

In early tumorigenesis, immune cells, including CD8+ T cells, detect danger signals from cancer cells and respond by eliminating these cells, leading to tumour immunogenicity. Dendritic cells are key regulators of tumour-specific immune responses in that they function as antigen presenting cells, activating CD8+ cytotoxic T lymphocyte (CTL) responses or stimulating CD4+ T cells through interaction with MHC class II molecule–antigen complexes (Finn 2008). In pancreatic tumorigenesis, Hiraoka and colleagues reported that chemokine (C-X-C motif) ligand 17 (CXCL17) was responsible for infiltration of immature dendritic cells while the intercellular adhesion molecule (ICAM2) permitted killing of tumour cells by CD8+ cells. During evolution from precursor lesions to PDAC they found that the host immune response was tempered and tumours developed immune tolerance by down-regulating CXCL17 and ICAM2 (Hiraoka et al, 2011). Interestingly, Dillman and colleagues recently described the value of reintroducing dendritic cells loaded with antigen from autologous irradiated tumour cells in metastatic melanoma. Those patients who received this dendritic cell vaccine survived significantly longer than those irradiated with tumour cell vaccine alone (Dillman et al, 2011). Conversely, in PDAC, blockade of Toll-like receptor 4 (TLR4) signalling enhanced dendritic cell-mediated recruitment of T helper 2 (Th2) CD4+ cells, and this resulted in increased pancreatic inflammation and accelerated tumour progression (Ochi et al, 2012). Thus, depending on antigenic stimulus, dendritic cells can both promote and inhibit tumour progression.

Once tumour immune surveillance is overcome, the composition of the immune infiltrate changes and a pro-tumorigenic leukocyte profile emerges (Figure 1). It is not yet fully understood how these different pro and anti-tumoural components of the immune system are engaged, however, signals in the tumour microenvironment, including IL-10, can prevent dendritic cell activation, thus dampening the adaptive immune response (Koido et al, 2010). T cells can be both tumour suppressing and promoting depending upon their downstream target cells. CD4+ cells, particularly T regulatory cells and Th2 cells, increase in number over the course of pancreatic cancer progression, whereas CTL cells decrease (Clark et al, 2007). This dynamic circumstance leads to a progressive accumulation of immunosuppressive cells that inhibit the anti-tumoural immune response. In tumours infiltrated by T regulatory cells both innate and adaptive immunity may well be suppressed, while myeloid-derived suppressor cells (MDSCs) can also subdue anti-tumour activity (Gabrilovich and Nagaraj, 2009).

In this milieu of tumour cell killing and escape, the immune system must clear damaged tissue. Neutrophils move into tissues under the control of chemokine signalling, particularly CXCR2. Though short lived, neutrophils have the potential to induce significant tissue remodelling. Fridlender et al (2009) demonstrated that following transforming growth factor-β (TGF-β) blockade,
neutrophils exhibited an anti-tumour phenotype termed N1. Both neutrophils and MDSCs are Gr1⁺, CD11b⁺ cells. The progressive accumulation of MDSCs in PDAC is widely recognised, while their immunosuppressive effect is clear. They are known to have the capacity to differentiate into both neutrophils and macrophages (Kusmartsev et al, 2005), and it has been suggested that pro-tumorigenic N2 neutrophils (TANs) may differentiate from MDSCs of splenic origin. Indeed two groups have demonstrated that in response to oncogenic Kras-dependent GM-CSF secretion, MDSCs are recruited to the tumour micro-environment where they have a role in suppressing CTL cell action and thus permitting tumorigenesis (Bayne et al, 2012; Pylayeva-Gupta et al, 2012). When GM-CSF was suppressed, MDSCs failed to infiltrate the tumour microenvironment, and tumour growth was limited by infiltrating CTL cells. Furthermore, this effect was fully rescued by CTL depletion. Activated neutrophils also have the capacity to cause tissue damage via the release of matrix metalloproteinases (MMPs), while production of reactive oxygen species may promote mutagenesis. Reactive oxygen species have been implicated in the activation of NF-κB, which has been intimately associated with the inflammatory response and PDAC progression (DeNicola et al, 2011).

Tumour-associated macrophages (TAMs) are present in early PDAC and persist throughout progression, representing an important component of the PDAC-associated leukocyte infiltrate (Clark et al, 2007). Intriguingly, TAMs can evolve to exhibit an M2 phenotype that is pro-tumorigenic in terms of promotion of growth and angiogenesis, and suppression of adaptive immunity (Sica and Mantovani, 2012). It is thought that this polarisation of macrophages from a tumour-suppressive M1 phenotype is initiated by cytokine signals, including IL-10 and TGF-β, received from T regulatory and tumour cells (Sica and Mantovani, 2012). This situation can be thought of as self-reinforcing, as production of pro-angiogenic molecules, including vascular endothelial growth factor (VEGF), attracts yet more macrophages to the tumour microenvironment, further enhancing angiogenesis. The plasticity of TAMs makes them an ideal therapeutic target for manipulation to generate an anti-tumour response.

Understanding the regulation of cancer immunosurveillance may help efforts to promote adaptive immune responses against PDAC, whereas the plasticity of TAMs and TANs may provide therapeutic opportunities for the promotion of an anti-tumour phenotype. There are examples where TAMs have been seen to elicit cancer-destructive properties. Hagemann et al (2008) have suggested that macrophages may be re-educated towards an anti-tumour function through control of NF-κB activity. Triggering a ‘good’ innate immune response is a potential goal for anti-cancer therapy with the activation of M1 macrophages and inhibition of MDSCs and T regulatory cells as the important objectives.

Antibodies against M2 macrophages and regulatory T cells are now available for trial in preclinical models, while measures to harness the immune system to eradicate tumour cells are being considered. An excellent example was recently provided in a cohort of metastatic PDAC patients who received CD40-targeting monoclonal antibodies in addition to gemcitabine (Beatty et al, 2011). CD40 activation, somewhat unexpectedly, led to macrophage-dependent tumour regression. Promotion of properties of the adaptive immune system that protect us from malignancy in health holds promise for future trials.

THE IMPORTANCE OF INFLAMMATION IN PDAC DEVELOPMENT

Mutations in the KRAS oncogene are presumed responsible for PDAC initiation and seen in the earliest PanIN lesions in nearly all cases. As these lesions progress from PanIN2 to PanIN3, mutations in the tumour suppressors TP53, CDKN2A, DPC4/SMAD4 and BRCA2 are observed at varying frequencies, leading to invasive PDAC (HRuban et al, 2000). Recently, attention has switched to the elucidation of the role played by inflammation in the evolution of PDAC. Mouse models have been useful in demonstrating the importance of local inflammation in pancreatic tumorigenesis. The Barbacid lab has shown that experimentally induced chronic pancreatitis (CP) combined with targeted expression of oncogenic mutant KrasG12D to pancreatic acini, cooperate to induce PDAC formation (Guerra et al, 2007). Lee and Bar-Sagi (2010)...

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demonstrated that on a background of inflammation, rather than activating senescence, \textit{Kras} mutation led to decreased expression of \textit{Ink4a} and a block in senescence, leading the authors to conclude that mutation of \textit{Kras} in the presence of a pre-existing inflammation would lead rapidly to tumorigenesis. These studies suggest that PDAC can occur on a background of pre-existing, possibly subclinical, pancreatic inflammation. The identification of patients at risk of low-level pancreatic inflammation could allow clinical application of these findings in future.

Interestingly, patients with CP have an increased risk of PDAC, though the majority of patients do not develop cancer (Hezel \textit{et al.}, 2006). Patients presenting to clinics with CP represent the extreme clinical application of these findings in future. Patients at risk of low-level pancreatic inflammation could allow possibly subclinical, pancreatic inflammation. The identification of inflammation would lead rapidly to tumorigenesis. These studies suggest that PDAC can occur on a background of pre-existing, inflammation which would rapidly to tumorigenesis. These studies suggest that PDAC can occur on a background of pre-existing, inflammation would lead rapidly to tumorigenesis.

The mounting of a response to tumour threat or injury is not the only way in which the immune system can become pathologically activated in PDAC. In chronic inflammation-associated PDAC, NF-κB expression leads to secretion of the pro-tumorigenic Shh ligand by macrophages (Yamasaki \textit{et al.}, 2010). When stimulated by cytokines such as TNF-α, NF-κB activates expression of growth-promoting molecules including cyclin D1 and E, c-Myc and IL-6. There is also evidence that mutant \textit{Kras} can activate NF-κB through IL-1α and p62. Meilisi \textit{et al.} identified autocrine IL-1α secretion from PDAC cells to be the mechanism of constitutive NF-κB activation. They showed that in response to IL-1α stimulation, PDAC cells exhibited an invasive phenotype \textit{in vitro} and when injected orthotopically into the murine pancreas generated liver metastases. This phenotype was reversed upon inhibition of NF-κB (Meilisi \textit{et al.}, 2009). Interestingly IKK2/β, a kinase required to release NF-κB from inhibition, is necessary for the induction of CP, and PanIN and PDAC formation following \textit{Kras} mutation (Ling, 2012). When genetically inactivated in a normally rapid mouse model of PDAC driven by mutant \textit{Kras} and conditional deletion of \textit{Cdkn2a}, mice fail to develop PDAC over a 1-year period. In addition, proliferation was significantly reduced in \textit{Pdx1-Cre; KrasG12D+/−; IKK2/β−/−} animals compared with \textit{Pdx1-Cre; KrasG12D+/−} controls. Further, the inflammatory infiltrate seen in \textit{Pdx1-Cre; KrasG12D+/−; IKK2/β−/−} animals is lower than that seen in \textit{Pdx1-Cre; KrasG12D+/−; IKK2/β−/−} controls, which included T and B cells, macrophages and neutrophils, which was not observed in \textit{Pdx1-Cre; KrasG12D+/−; IKK2/β−/−} animals. These observations provide further evidence supporting a role for inflammation in PDAC development and a critical role for NF-κB in this process.

Thus, murine models have provided compelling evidence for the dependence of PDAC initiation on intrinsically generated

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**PDAC-ASSOCIATED MUTATIONS SUSTAIN AN INFLAMMATORY MICROENVIRONMENT**

The mounting of a response to tumour threat or injury is not the only way in which the immune system can become pathologically activated in PDAC. Initiating \textit{Kras} mutations induce upregulation of inflammatory pathways in PDAC, and downstream of \textit{Kras} the RAF signalling pathway induces production of chemokines and cytokines that may in themselves be pro-tumorigenic, including chemokines CXCL1 and CXCL8, the major ligands for CXCR2 (Balkwill, 2012). The oncogenic transcription factor c-Myc, also activated in tumours, promotes cell-autonomous proliferation and is key to tumour microenvironment remodelling via stimulation of inflammatory cells and cytokine production (Balkwill, 2012).

Ochi \textit{et al.} recently showed high expression of Toll-like receptor 7 (TLR7) in PDAC. They elegantly demonstrated in a murine model that TLR7 ligation accelerated pancreatic tumorigenesis via upregulation of inflammatory cascades including STAT3 and NF-κB pathways. Toll-like receptor 7 blockade protected against carcinogenesis and when mice lacking TLR7 exclusively within their inflammatory cells were generated, they were completely protected from carcinogenesis (Ochi \textit{et al.}, 2012). Toll-like receptor 7 inhibition may have a role in combating the tumour microenvironment in PDAC by preventing signalling through downstream inflammatory cascades.

A number of upregulated inflammatory and oncogenic pathways in PDAC converge on the transcription factors STAT3 and NF-κB, suggesting that these are excellent therapeutic targets. Two recent studies have demonstrated upregulation of STAT3 in PDAC (Fukuda \textit{et al.}, 2011; Lesina \textit{et al.}, 2011). Using mouse models of endogenous mutant \textit{Kras} plus experimentally induced pancreatitis, the authors observed STAT3 activation in response to acinar damage. Activation was only sustained in the setting of \textit{Kras}-accelerated metaplastic ducts and PanIN formation. When STAT3 was deleted, specifically within the pancreas, this significantly interfered with inflammation-mediated PanIN formation, and reduced both inflammatory cell infiltration (CD45+ cells) and cytokine expression including IL-6. Candidate ligands for activation of STAT3 signalling in these model tumours were interrogated, and IL-6 was significantly upregulated and expressed predominantly by infiltrating macrophages. In human PDAC, serum IL-6 levels were significantly raised and suggested a potential link between systemic inflammation and PDAC tumorigenesis. Importantly IL-6 activation can occur via trans-signalling when IL-6 binds to a naturally occurring soluble form of IL-6R and forms a complex that induces IL-6-specific signalling in cells that lack a membrane-bound receptor. Indeed this latter form of signalling was found to be critical in the process of PDAC tumorigenesis. Furthermore Corcoran \textit{et al.} (2011) have established that pSTAT3 levels predict PDAC cell sensitivity to JAK2 inhibition, suggesting pSTAT3 status may be used in future to select patients for JAK/STAT inhibition.

Several studies have shown NF-κB signalling to be an important signalling node in pancreatic cancer. In chronic inflammation-associated PDAC, NF-κB expression leads to secretion of the pro-tumorigenic Shh ligand by macrophages (Yamasaki \textit{et al.}, 2010).

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inflammation. The activation of NF-κB and STAT3 signalling by upregulated systemic cytokines including IL-6, IL-1α, and TNF-α identifies many of these molecules as excellent therapeutic targets. Anti-TNF-α therapies such as infliximab are well established in inflammatory conditions such as inflammatory bowel disease. Egberts et al (2008) demonstrated in orthotopic PDAC models that anti-TNF therapy reduced primary tumour size and metastases, however, these results have not been translated to clinical trials. Targeting IL-6, a recognised mediator of PDAC tumorigenesis, may be of therapeutic benefit. IL-6 monoclonal antibodies, Siltuximab and Tocilizumab, which bind to the soluble form of the IL-6 receptor are available for trial and currently under assessment in ovarian cancer (clinicaltrials.gov). Furthermore, key nodes in the transmission of inflammatory signals such as STAT3, which is strongly implicated in PDAC tumorigenesis, can now be targeted with direct inhibitors. Recently triterpenoids and rexinoids demonstrated good efficacy in STAT3 binding, and prolonged survival in the KrasG12D; p53fl/+ mouse model (Liby et al, 2010). These studies suggest that the repositioning of some anti-inflammatory to treat pancreatic cancer may offer clinical benefit in the future.

### COOPERATION BETWEEN TSG MUTATION AND CHEMOKINE SIGNALLING IN PDAC PROGRESSION

Recent studies support the deregulation of CXC chemokines in late-stage PDAC. Mutation of p53 can drive migration and metastasis in PDAC and this process was recently shown to be dependent on NF-κB signalling downstream of CXCR2 ligand upregulation (Li et al, 2011). In addition, CXCL5, a ligand for CXCR2, was overexpressed in human PDAC and marked patients with poor outcome, advanced tumour stage and increased tumour size. Chemokine signalling may be critical in promoting invasion and metastasis in PDAC, especially downstream of mutations in TP53.

Chemokine signalling may also be deregulated by SMAD4 mutations, found in 50% of PDAC cases (Rozenblum et al, 1997). SMAD4 is a TGF-β signalling molecule, and its loss is thought to influence PDAC tumorigenesis through coordination of TGF-β signalling in the tumour and stroma, as both fibroblasts and epithelial cells respond to TGF-β in the tumour microenvironment (Ijichi et al, 2011). In a mutant Kras mouse model, Ijichi et al (2011) demonstrated that TGF-β receptor II knockout resulted in aggressive PDAC that histologically recapitulated human disease. Secretion of CXCR2 specific chemokines, including CXCL1 and CXCL5, was significantly increased in these mice, and regulated by TGF-β signalling and NF-κB. Interestingly, stromal fibroblasts also expressed significantly higher CXCR2 levels compared with epithelial cells. Treatment with a CXCR2 inhibitor over a short period improved survival and reduced microvessel density, supporting the theory that CXCR2 ligands are important in driving tumour progression following TSG mutation in PDAC.

These studies support the notion that pro-inflammatory pathways can be regulated by multiple different driver mutations in PDAC. The prognostic significance of deregulated chemokines and the functional role they have following TSG mutation, together...
with the successful use of their inhibitors in other diseases, suggest that chemokines are potentially exciting therapeutic targets in patients with PDAC.

**METASTASIS AND INFLAMMATION**

Recent studies have shown metastases may occur in PDAC even before a primary tumour has formed, a behaviour associated with early epidermal mesenchymal transformation (Rhim et al., 2012). This process was accelerated in the presence of pancreatic inflammation, while the most invasive areas of tumour were seen at foci of inflammation. This phenotype was suppressed by dexamethasone, highlighting the integral role played by tumoural inflammation. CXCR2 has been shown to be an important pro-survival factor in metastatic breast cancer models (Acharyya et al., 2012). CXCR2 inhibition in this setting sensitised metastases to killing by standard chemotherapeutics. Chemokine-specific small molecules and monoclonal antibodies are undergoing preclinical study at present. The identification of inflammatory signalling pathways in the generation and maintenance of metastases in PDAC provides hope for combinatory anti-metastatic therapies in the near future.

**CONCLUSIONS**

Treating PDAC by targeting its reliance on the tumour microenvironment may be approached from three aspects (Figure 2): first by promotion of-tumour-eliminating processes; second by suppression of tumour-promoting inflammation; and third via modulation of the protective fibrotic stroma of PDAC to allow access to tumour epithelium by conventional chemotherapeutics. The availability of preclinical in vivo models that accurately recapitulate the histological progression of PDAC and its complex microenvironment provides an excellent tool linking basic science with clinical application. The potential for preoperative characterisation of PDAC and direction of tumour-specific individualised therapy is within sight. Individualised multi-targeted therapy is likely to be necessary in order to treat PDAC effectively, as high recurrence rates following surgery and late presentation of disease remain significant hurdles.

Only 10–15% of PDAC are resectable at presentation highlighting that one of the greatest challenges facing PDAC management is earlier diagnosis. However, even those with R0 resection margins following surgery sadly suffer disease recurrence despite adjuvant therapy, often due to acquired drug resistance.

We have outlined the importance of the activated tumour microenvironment in PDAC maintenance and progression. Inflammation is implicated in the earliest stages of PDAC tumorigenesis, is strongly implicated in early metastases, and in tumour progression following TSG mutation. Combinatorial therapeutic regimens must look to capitalise on the importance of this tumour–microenvironment relationship. Aggressive targeted neoadjuvant therapy in surgically resectable patients should be considered, while those with more advanced disease may well respond to similar regimens. Incorporating newly developed anti-inflammatory agents alongside standard chemotherapeutics may help provide much needed improvement in tumour sensitivity to therapy for patients suffering PDAC.

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