

# Original research

# Chemotherapy-induced infiltration of neutrophils promotes pancreatic cancer metastasis via Gas6/AXL signalling axis

Gaia Bellomo,<sup>1</sup> Carolyn Rainer,<sup>1</sup> Valeria Quaranta,<sup>1</sup> Yuliana Astuti,<sup>1</sup> Meirion Raymant,<sup>1</sup> Elzbieta Boyd,<sup>1</sup> Ruth Stafferton,<sup>1</sup> Fiona Campbell,<sup>1</sup> Paula Ghaneh,<sup>1</sup> Christopher M Halloran,<sup>1</sup> Dean E Hammond,<sup>2</sup> Jennifer P Morton,<sup>3,4</sup> Daniel Palmer,<sup>1</sup> Dale Vimalachandran,<sup>1</sup> Robert Jones,<sup>1</sup> Ainhoa Mielgo,<sup>1</sup> Michael C Schmid <sup>1</sup>

Additional supplemental material is published online only. To view, please visit the journal online (http://dx.doi.org/ 10.1136/gutinl-2021-325272).

<sup>1</sup>Molecular and Clinical Cancer Medicine, University of Liverpool, Liverpool, UK <sup>2</sup>Molecular Physiology and Cell Signalling, University of Liverpool, Liverpool, UK <sup>3</sup>Cancer Research UK Beatson Institute, Glasgow, UK <sup>4</sup>Institute of Cancer Sciences, University of Glasgow, Glasgow, UK

#### Correspondence to

Professor Michael C Schmid, Molecular and Clinical Cancer Medicine, University of Liverpool, Liverpool, UK; mschmid@liverpool.ac.uk

GB and CR contributed equally.

Received 28 May 2021 Accepted 19 December 2021 Published Online First 12 January 2022



© Author(s) (or their employer(s)) 2022. Re-use permitted under CC BY. Published by BMJ.

To cite: Bellomo G, Rainer C, Quaranta V, et al. Gut 2022;71:2284-2299

ABSTRACT **Objective** Pancreatic ductal adenocarcinoma (PDAC) is a highly metastatic disease and cytotoxic chemotherapy is the standard of care treatment for patients with advanced disease. Here, we investigate how the microenvironment in PDAC liver metastases reacts to chemotherapy and its role in metastatic disease progression post-treatment, an area which is poorly understood.

**Design** The impact of chemotherapy on metastatic disease progression and immune cell infiltrates was characterised using flow and mass cytometry combined with transcriptional and histopathological analysis in experimental PDAC liver metastases mouse models. Findings were validated in patient derived liver metastases and in an autochthonous PDAC mouse model. Human and murine primary cell cocultures and ex vivo patient-derived liver explants were deployed to gain mechanistical insights on whether and how chemotherapy affects the metastatic tumour microenvironment.

**Results** We show that in vivo, chemotherapy induces an initial infiltration of proinflammatory macrophages into the liver and activates cytotoxic T cells, leading only to a temporary restraining of metastatic disease progression. However, after stopping treatment, neutrophils are recruited to the metastatic liver via CXCL1 and 2 secretion by metastatic tumour cells. These neutrophils express growth arrest specific 6 (Gas6) which leads to AXL receptor activation on tumour cells enabling their regrowth. Disruption of neutrophil infiltration or inhibition of the Gas6/AXL signalling axis in combination with chemotherapy inhibits metastatic growth. Chemotherapy increases Gas6 expression in circulating neutrophils from patients with metastatic pancreatic cancer and recombinant Gas6 is sufficient to promote tumour cell proliferation ex vivo, in patient-derived metastatic liver explants.

Conclusion Combining chemotherapy with Gas6/ AXL or neutrophil targeted therapy could provide a therapeutic benefit for patients with metastatic pancreatic cancer.

### BACKGROUND

Metastasis is the leading cause of cancer-related death. Pancreatic ductal adenocarcinoma (PDAC) frequently metastasizes to the liver<sup>1 2</sup> and liver metastasis is accompanied by the formation of an

### Significance of this study

#### What is already known on this subject?

- $\Rightarrow$  Pancreatic cancer is a devastating metastatic disease for which better therapies are urgently needed.
- $\Rightarrow$  Pancreatic cancer frequently metastasises to the liver where the metastatic microenvironment facilitates the seeding and growth of metastases.
- $\Rightarrow$  Cvtotoxic chemotherapy is the standard care of treatment for all patients with pancreatic cancer, including those with locally advanced or metastatic disease, and as adjuvant treatment for patients after surgical resection of their primary tumour.
- $\Rightarrow$  It is unclear how the metastatic microenvironment reacts to chemotherapy and its role in metastatic disease progression post-treatment.

### What are the new findings?

- $\Rightarrow$  Cessation of chemotherapy induces the recruitment of neutrophils to the liver, resulting in increased metastatic growth.
- $\Rightarrow$  Neutrophils are recruited to the liver via CXCL1 and 2 expression by disseminated pancreatic cancer cells.
- $\Rightarrow$  Neutrophils recruited to the liver postchemotherapy express growth arrest specific 6 (Gas6) which leads to AXL receptor activation on tumour cells.
- $\Rightarrow$  Gas6-mediated activation of the AXL receptor on tumour cells promotes the regrowth of tumour cells after chemotherapy treatment in vitro and in vivo.
- $\Rightarrow$  Disruption of neutrophil infiltration or inhibition of the Gas6/AXL signalling axis in combination with chemotherapy inhibits metastatic growth.

inflammatory-fibrotic metastatic microenvironment that supports the colonisation and outgrowth of disseminated cancer cells.<sup>3-6</sup> Myeloid immune cells, including monocytes, macrophages and neutrophils, are found in high numbers in the metastatic niche and have been shown to promote the metastatic process.<sup>7-9</sup> Macrophages are highly plastic cells and, depending on their activation state, can acquire tumour supportive or tumour repressive



# How might it impact on clinical practice in the foreseeable future?

⇒ Combining chemotherapy with Gas6/AXL or neutrophil targeted therapy may offer a new opportunity in the treatment of patients with metastatic pancreatic cancer and in the adjuvant setting of patients that have undergone tumour resection.

functions.<sup>10 11</sup> During liver metastasis, macrophages are prometastatic, display an immunosuppressive phenotype,<sup>12 13</sup> and promote fibrosis.<sup>3</sup> Emerging evidence suggests that neutrophils play a critical role during the early steps of metastasis.<sup>14</sup> Neutrophils can promote the colonisation of the distant site through the release of neutrophil extracellular traps (NETs),<sup>15 16</sup> induction of angiogenesis,<sup>9 17</sup> secretion of leukotrienes<sup>18</sup> and by their immunosuppressive activities.<sup>19 20</sup> However, whether myeloid immune cell functions in pancreatic cancer metastases are altered in response to therapeutic interventions remains unknown.

Systemic spread is an early event in pancreatic cancer progression<sup>1</sup> and by the time PDAC patients are diagnosed, the majority (~80%) present with non-resectable metastatic cancer.<sup>2</sup> A total of 15%–20% of PDAC patients are eligible for surgical resection of their primary tumour. However, clinically undetectable micrometastatic lesions are often already present at the time the primary tumour is removed, and more than 70% relapse with distant metastasis within 24 months of surgery.<sup>21</sup> The time of recurrence after surgical resection strongly correlates with overall survival, and an early hepatic metastatic relapse is associated with the worse prognosis.<sup>22</sup> Following diagnosis of liver metastases, median survival on systemic chemotherapy is just 9 months.<sup>23</sup>

Cytotoxic chemotherapy is the standard care of treatment for all patients with pancreatic cancer, including those with locally advanced or metastatic disease and as adjuvant treatment for patients after surgical resection of their primary tumour.<sup>24</sup> Gemcitabine, gemcitabine/capecitabine, nab-paclitaxel and FOLFIRINOX are the most common chemotherapeutic treatment options.<sup>25</sup> Although the effect of chemotherapy on the primary tumour site is well characterised,<sup>21</sup> our understanding of how chemotherapy shapes the hepatic metastatic microenvironment and how this affects metastatic disease progression remains unknown. A better understanding of this process could lead to treatments that improve the efficacy of current systemic chemotherapies.

#### RESULTS

# Gemcitabine treatment restrains metastatic progression, but disease relapses when treatment is withdrawn

To model chemotherapeutic treatments of metastatic pancreatic cancer in vivo, we induced PDAC liver metastasis in mice by intrasplenic implantation of KPC derived cells and initiated gemcitabine treatment once metastatic lesions had been established (at day 12 postimplantation),<sup>3</sup> (figure 1A). While KPC cancer cells were sensitive to gemcitabine when treated in vitro (online supplemental figure S1A), gemcitabine treatment did not improve the overall survival of animals with pancreatic cancer liver metastasis (figure 1A). Bioluminescent in vivo imaging analysis revealed that metastatic tumour burden was significantly reduced in the gemcitabine treated animals at the end of the treatment schedule (d22) (figure 1B), but no differences in tumour

burden were detected at the humane endpoints (between day 32 and day 48 (figure 1C). H&E staining of liver tissue sections further confirmed a significant reduction of metastatic tumour lesions by the end of the treatment schedule (day 22), while this reduction was no longer detected at humane endpoints (online supplemental figure S1B-E). Postmortem analysis proofed extensive tumour burden in the liver, while tumour formation in the spleen remained minor (online supplement figure S1F, G). We next assessed tumour cell death in livers from control (saline treated) versus gemcitabine treated animals. We found that the percentage of apoptotic cancer cells, assessed by cleaved caspase 3 (CC3) staining, was significantly increased in gemcitabine treated animals compared with control tumour-bearing mice by the end of the treatment schedule (day 22) (figure 1D,E). However, by humane endpoints, after withdrawal of gemcitabine treatment, the initially observed increase in cancer cell death was lost (online supplemental figure S1H, I). Adjuvant chemotherapy is the established standard-of-care for patients who undergo surgical resection of their primary pancreatic tumour.<sup>24</sup> The short time and the high frequency at which these patients relapse with metastatic disease (median of 9 months after resection<sup>23</sup>) strongly suggests that occult micrometastases were already established at the time of surgery.<sup>26</sup> To test the effect of chemotherapy on micrometastatic lesions equivalent to the adjuvant treatment setting, we next administered a single dose of gemcitabine at day 3 post-tumour implantation, after the initial seeding period and where micro-metastatic lesions are present (figure 1F).<sup>3 13</sup> Similar to what we observed with larger metastatic lesions, administration of gemcitabine also reduced micrometastatic tumour burden (figure 1G,H) and tumour lesion areas in the liver at day 4 (figure 1I, J), but this effect was lost at day 14 (figure 1G–J). By 24 hours after gemcitabine administration, the percentage of apoptotic cancer cells (TUNEL+) in micrometastatic lesions was markedly increased in gemcitabine treated animals compared with control animals (figure 1K,L). However, at day 14, the rate of TUNEL +cancer cells declined in the gemcitabine treated tumours to similar levels as in the untreated cohort (online supplemental figure S1J, K). Macrophages and dendritic cells (DCs) are phagocytic cells from the innate arm of the immune system that play a key role in the removal of dead cells and are critical for the induction of an antitumour immune response in cancer.<sup>27</sup> In order to identify the phagocytosis of cellular cancer debris by macrophages and DCs, we used flow cytometry to measure the fluorescent signal of zsGreen labelled cancer cells within macrophages (CD11b<sup>+</sup>F4/80<sup>+</sup>CD11c<sup>neg</sup>), CD11b<sup>+</sup> DCs (CD11c<sup>+</sup>CD11b<sup>+</sup>CD103<sup>neg</sup>F4/80<sup>neg</sup>) and CD103<sup>+</sup> DCs (CD11c<sup>+</sup>CD103<sup>+</sup>CD11b<sup>neg</sup>F4/80<sup>neg</sup>). All three cell populations showed a significant uptake of zsGreen signal after gemcitabine administration compared with the control group (figure 1M; online supplemental figure S1L), suggesting an activation of innate immune cells in metastatic lesions in response to chemotherapy. Taken together, these findings show that gemcitabine induces cancer cell death in PDAC metastatic lesions, but tumour growth relapses after treatment withdrawal and overall survival remains unchanged.

#### Gemcitabine treatment induces a short-term activation of a proinflammatory immune response in metastatic hepatic lesions

Since chemotherapy can promote the activation of an immune response in cancer<sup>27</sup> we next investigated, in more detail, the immune cell activation on gene expression level in metastatic lesions during the initial response to gencitabine treatment (day



**Figure 1** Gemcitabine restrains metastatic progression during treatment, but disease relapses and overall survival remain unchanged when treatment is withdrawn. (A–E) Liver metastasis was induced by intrasplenic implantation of  $1 \times 10^{6}$  KPC<sup>luc/zsGreen</sup> cells. Starting day 12, animals were treated with gemcitabine (100 mg/kg) or control (vehicle) every 3 days with four doses in total. (A) Survival analysis of gemcitabine and control-treated mice-bearing liver metastasis; log-rank (Mantel-Cox) test, p=0.2499. Median survival for control was 22 days (n=6 mice) and gemcitabine 33.5 days (n=8 mice) after treatment initiation. (B) Representative images of bioluminescence imaging (BLI) taken 1 day after last treatment dose (day 22). (C) Tumour burden assessed by BLI in gemcitabine treated group (n=8 mice) compared with control group (n=6 mice) at day 22 and humane endpoint (HEP). (D, E) Representative immunofluorescent images (D) and quantification (E) of apoptotic KPC<sup>luc/zsGreen</sup> cells staining positive for cleaved caspase 3 (CC3) at day 22 (n=5 mice /group). White arrowheads indicate apoptotic (CC3+) cancer cells. (F–L) Liver metastasis was induced by intrasplenic implantation of  $5 \times 10^5$  KPC<sup>luc/zsGreen</sup> cells and animals received one dose of gemcitabine (100 mg/kg) or control (vehicle) at day 3 (F). (G, H) Representative BLI images of dissected livers (G) and change in tumour burden (H) (day 4: n=5 mice/group/time point). (I, J)Representative images of H&E-stained liver sections (I) and quantification (L). White arrowheads indicate apoptotic (TUNEL+) cancer cells. (M) uptake of apoptotic zsGreen-labelled KPC FC1199<sup>luc/zsGreen</sup> cancer cells by dendritic cells (DC) and macrophages (MACS) was evaluated 1 day after gemcitabine treatment. Frequency of zsGreen +cells among CD103<sup>+</sup> DC, CD11b<sup>+</sup> DC and MACS (n=5 mice/group). Scale bar 50 µM. Data are presented as mean±SEM. Unpaired t-test was used to calculate p values. \*P<0.05; \*\*p<0.01. H, healthy liver; M, metastase; n.s., not significant.

4) and after withdrawal (day 14) using the Mouse PanCancer Immune Profiling Panel (NanoString Technologies). Hierarchical clustering of the generated pathway scores revealed that gemcitabine induces distinct transcriptional changes during the initial response, as highlighted by the separate clustering of the gemcitabine groups compared with control groups (figure 2A, left). However, the distinct signatures between control and gemcitabine-treated metastatic lesions were lost after withdrawal, as indicated by the loss of segregation between the two groups (figure 2A, right). Among the top upregulated pathways, we identified innate immune activation and T cell functions which are characteristic of an antitumour immune response (figure 2B). However, after gemcitabine withdrawal, these immune stimulatory pathways were markedly downregulated, suggesting that gemcitabine only triggers a temporal activation of an antitumour immune response in tumour-bearing mice (figure 2C). We next analysed disaggregated metastatic lesions by mass and flow cytometry to assess immune cell infiltration and their activation state. We found that during the initial response, macrophage numbers (CD45<sup>+</sup>CD11b<sup>+</sup>Ly6G<sup>neg</sup>F4/80<sup>+</sup>) significantly increased, and inflammatory monocytes numbers (CD45<sup>+</sup>CD-11b<sup>+</sup>Ly6C<sup>high</sup>Ly6G<sup>neg</sup>F4/80<sup>neg</sup>) were reduced (figure 2D; online supplemental figure S2A, B). In addition, CD4<sup>+</sup> T cell numbers significantly increased in response to treatment (online supplemental figure S3A, C). However, after chemotherapy withdrawal, neutrophils (CD45+CD11b+F4/80negLy6G+) and patrolling monocytes (pMo; CD45+CD11b+Ly6C<sup>low</sup>F4/80<sup>low/</sup> <sup>neg</sup>MHCII<sup>neg</sup>) increased the most in gemcitabine-treated tumours compared with control treated tumours, while T cell numbers were significantly decreased (figure 2E). The decrease in total T cell numbers was most likely due to a reduction in CD8<sup>+</sup> T cells, since the less abundant CD4<sup>+</sup>FoxP3<sup>+</sup> T regulatory cells (T<sub>res</sub>) rather increased (online supplemental figure S3A-D). Consistent with an antitumour immune response, we found a significant increase in the activation of CD8<sup>+</sup> and CD4<sup>+</sup> T cells (figure 2F; online supplemental figure S3E, F), DCs (figure 2G), macrophages (figure 2H) and NK cells (figure 2I) during the initial response to gemcitabine (online supplemental figure S2A, B). Again, this effect was lost after withdrawal of the treatment.

Further analysis of metastatic liver tissues confirmed that gemcitabine treatment induces the overall accumulation of macrophages (F4/80<sup>+</sup>), particularly of macrophages with a proinflammatory phenotype (iNOS<sup>+</sup>) (figure 2J) while macrophages with an immunosuppressive phenotype (CD206<sup>+</sup> and YM1<sup>+</sup>) were reduced (online supplemental figure S3G-I). However, no significant changes in macrophage infiltration or activation were observed after treatment withdrawal (figure 2K; online supplemental figure S3K, L). Taken together these data suggest that gemcitabine administration induces the activation of an antitumourigenic immune response at the metastatic site, characterised by an increase in proinflammatory macrophages, activated CD8<sup>+</sup> T cells and NK cells. However, after treatment withdrawal the initial immune cell activation is lost and metastatic lesions revert back to an immunosuppressive microenvironment, which is commonly found in established metastatic PDAC tumours.<sup>12 13</sup>

#### Macrophage depletion after gemcitabine treatment increases CD8+ T cell infiltration, but neutrophil depletion has no effect on CD8+ T cell numbers

Since neutrophils and macrophages can both effectively suppress  $CD8^+$  T cell responses,<sup>11 14</sup> we next questioned whether the depletion of either of these myeloid cell types is sufficient to stop

metastatic relapse and to sustain the initially observed CD8<sup>+</sup> T cell response (figure 2B and F). To address this question, we ran two separate depletion studies using monoclonal antibodies targeting neutrophils (aLy6G) or macrophages (aCSF-1) in the presence or absence of gemcitabine treatment. Liver metastasis was induced by intrasplenic implantation of KPC cells. After 3 days, mice-bearing micrometastatic lesions were treated with gemcitabine or saline (control) and 1 day later (day 4) we commenced the depletion of neutrophils (figure 3A-C) or macrophages (figure 3D-F) using  $\alpha$ Ly6G and  $\alpha$ CSF-1 antibodies, or their corresponding isotype controls (IgG). Depletion of neutrophils after gemcitabine administration markedly reduced the metastatic tumour burden compared with gemcitabine/IgG treatment (figure 3B,C, online supplemental figure S4A, B), but depletion of neutrophils in the absence of gemcitabine did not have any effect on metastatic tumour burden (online supplemental figure S4C). In contrast, depletion of macrophages by  $\alpha$ CSF-1 significantly reduced metastatic tumour burden in both saline (control) and gemcitabine treated mice (figure 3E,F; online supplemental figure S4D, E). Flow cytometry analysis of disaggregated metastatic lesions derived from gemcitabine treated animals revealed that the depletion of macrophages increased CD8<sup>+</sup> T cell infiltration, while neutrophil depletion did not affect CD8<sup>+</sup> T cell infiltration (figure 3G). In agreement with these findings, we did not detect an increase in Granzyme B expression in CD8<sup>+</sup> T cells in gemcitabine-treated animals where neutrophils were depleted (figure 3H), but we found a significant increase of Granzyme B expressing CD8<sup>+</sup> T cells in metastatic lesions of gemcitabinetreated mice where macrophages were depleted (figure 3J,K). We also confirmed that applied macrophage-depletion and neutrophil-depletion strategies indeed reduced their corresponding numbers at the metastatic site (online supplemental figure S4F-H). Notably, neutrophil-depletion or macrophagedepletion after gemcitabine treatment also increased overall survival of the mice compared with gemcitabine treatment alone (figure 3L,M). Taken together, these data show that gemcitabine administration is accompanied by an infiltration of macrophages during the initial response, while neutrophils are recruited to the metastatic site after therapy withdrawal. Depletion of macrophages or neutrophils after gemcitabine withdrawal enhances the therapeutic effect of gemcitabine. Notably, while macrophage depletion restores CD8<sup>+</sup> T cell infiltration and activation, neutrophil depletion does not affect CD8<sup>+</sup> T cells, suggesting that neutrophils promote metastatic relapse in a CD8<sup>+</sup> T cell independent manner.

## Chemotherapy withdrawal triggers the recruitment of Gas6expressing neutrophils to hepatic metastatic tumours

Next, we further explored the mechanism by which neutrophils promote metastatic relapse after chemotherapy withdrawal. Neutrophil depletion after gemcitabine treatment reduced metastases in the presence and absence of  $CD8^+T$  cells (figure 4A; online supplemental figure S4I, J), suggesting that the neutrophils can directly affect cancer cell regrowth. Hence, we next assessed cancer cell proliferation in tumour sections after gemcitabine withdrawal (day 14). After gemcitabine withdrawal, metastatic deposits showed a significant increase of proliferating (Ki67<sup>+</sup>) cancer cells compared with metastatic deposits from the saline treated control group (figure 4B,C). Importantly, the depletion of neutrophils only reduced cancer cell proliferation associated with gemcitabine withdrawal, and had no impact on cancer cell proliferation in saline (control) treated mice, suggesting a treatment induced growth promoting function



Figure 2 Gemcitabine administration induces a short-term activation of a proinflammatory immune response in metastatic pancreatic cancer. (A-K) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc</sup>/<sup>zsGreen</sup> cells and animals were treated with gencitabine (100 mg/kg) or control (vehicle) at day 3. Metastatic livers were resected at initial response (day 4) and after withdrawal of treatment (day 14) for transcriptional, mass cytometry and tissue staining analysis. (A) Heatmap depicting hierarchical clustering of pathway scores (n=3 mice/group/time point). (B–C) Graph depicting top pathway scores observed in (B) metastatic livers of gemcitabine treated animals compared with control animals during initial response (day 4) and in (C) metastatic livers after gemcitabine withdrawal (day 14) compared with the initial response (day 4). (D, E) Coloured viSNE maps with each colour representing one immune cell population assessed by mass cytometry and quantification of main immune cell types among control (CTR) and gemcitabine (GEM) treated liver metastases at day 4 (A) and day 14 (B), respectively (CTR D4 n=4 mice, GEM D4 n=4 mice; CTR d14 n=3 mice; GEM d14 n=4 mice). (F–I) Quantification of metastasis infiltrating immune cells and their activation state by mass cytometry at initial treatment response (day 4) and after treatment withdrawal (day 14). (F)Cytotoxic CD8+ T cell activation (CD69+), (G) dendritic cell (DC) activation (CD86 +MHCII<sup>high</sup>), (H) macrophage activation (CD86 +MHCII<sup>high</sup>) and (I) natural killer (NK) cell activation (CD69+) (CTR D4 n=4 mice, GEM D4 n=4 mice; CTR d14 n=3 mice; GEM d14 n=4 mice). (J, K) Representative immunofluorescent images and quantification of iNOS + and F4/80+ macrophages in liver tumours during initial response (n=4 mice/group) (D) and after gemcitabine withdrawal (E) (n=3 mice in CTR group; n=4 mice in GEM group). White arrowheads indicate iNOS + macrophages. Scale bar 50 µM; M=metastases, H=healthy liver. Data are presented as mean±SEM. \*P<0.05; \*\*p<0.01; n.s., not significant, by unpaired t-test. For multiple comparisons (D, E), one-way ANOVA coupled with Dunnett's post hoc testing was performed. ANOVA; analysis of variance.



**Figure 3** Macrophage depletion increases CD8 + T cell infiltration, but neutrophil depletion has no effect on CD8+ T cell numbers. (A–M) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc</sup>/<sup>zsGreen</sup> cells and animals were treated with gemcitabine (100 mg/kg) or control (vehicle) at day 3. (A–C, G–I) At day 4, mice were treated with IgG control (CTR) or αLy6G antibody. Schematic illustrating experiment (A). Change in tumour burden was guantified by in vivo BLI (n=3 mice/group). Representative images (B) and guantification (C). (D–G, J, K) At day 4, mice were treated with IgG control (CTR) or  $\alpha$ CSF-1 antibody. Schematic illustrating experiment (D). Change in tumour burden was quantified by in vivo BLI (CTR n=3 mice;  $\alpha$ CSF-1 n=4 mice). Representative images (E) and guantification (F). (G) Change in CD8+T cell infiltration into metastatic lesions was guantified by flow cytometry analysis in mice treated with  $\alpha$ Ly6G or  $\alpha$ CSF-1 or their corresponding IgG controls. (H, I) Representative immunofluorescent images of CD8+GranzymeB + T cell staining of liver sections from mice treated with IgG or  $\alpha$ Ly6G (H) and quantification (I) of CD8+GranzymeB+T cells (GranzymeB=GzmB). (J, K) Representative immunofluorescent images of CD8+GranzymeB+T cell staining of liver sections from mice treated with IgG or αCSF-1 (J) and quantification (K) of CD8 +GranzymeB+ T cells. White arrowheads indicate CD8 +GranzymeB+ T cells. (L) Liver metastasis was induced by intrasplenic implantation of 1×10<sup>6</sup> KPC<sup>luc/zsGreen</sup> cells. At day 3, all animals were treated with gemcitabine (100 mg/kg). At day 4, mice were treated with IgG control (CTR) or  $\alpha$ Ly6G antibody. survival analysis of gemcitabine + IgG and gemcitabine +  $\alpha$ Ly6G antibody-treated mice-bearing liver metastasis; log-rank (Mantel-Cox) test, p=0.0022. Median survival for gemcitabine + IgG was 35 days (n=6 mice) and gemcitabine +  $\alpha$ Ly6G 48 days (n=6 mice). (M) same as (L), but at day 4, mice were treated with IgG control (CTR) or  $\alpha$ CSF-1R antibody. Survival analysis of gemcitabine + IgG and gemcitabine +  $\alpha$ CSF-1R antibody-treated mice-bearing liver metastasis; log-rank (Mantel-Cox) test, p=0.0168. Median survival for gemcitabine + IqG was 29.5 days (n=6 mice) and gemcitabine + αCSF-1R 45 days (n=6 mice). Scale bar 50 μM. Data are presented as mean±SEM. Unpaired t-test was used to calculate p values. \*P<0.05; \*\*p<0.01. H, healthy liver; M, metastases; n.s., not significant.



Figure 4 Neutrophils promote cancer cells proliferation and Gas6 is highly expressed by metastatic associated neutrophils after gemcitabine treatment. (A) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (GEM; 100 mg/kg) at day 3. At day 4, mice were treated with  $\alpha$ Ly6G or IgG controls for 2 weeks; at day 7, mice were treated with  $\alpha$ CD8 or IgG controls until end point (day 14). Change in tumour burden was guantified by ex vivo bioluminescence imaging (BLI) (n=5 mice/group). (B-C) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. Mice were treated with gemcitabine or saline 3 days postcell implantation. Treatment with  $\alpha$ Ly6G or control IgG started at D4 (n=4 mice/group). Livers were resected after 14 days and assessed by Ki67 staining (proliferation marker). Representative IHC images (B) and quantification of proliferating Ki67+ tumour cell frequency in metastatic livers (C). Inset: asterisks indicate ductal structures formed by metastatic tumour cells (red arrow head). (D)Colony formation assay of gemcitabine stressed KPC cells in the presence or absence of metastasis infiltrating neutrophils (+Lv6G) or macrophages (+F4/80) isolated from tumour-bearing livers of mice at day 14 after treatment with GEM or saline treated (CTR). Bar graph shows fold upregulation of BLI signal compared with Gem-treated KPC cells alone (red shaded) (three independent experiments; mean±SEM). (E) Quantification of Gas6 mRNA levels by real time PCR in intrametastatic pancreatic cancer cells, neutrophils (Ly6G), macrophages (F4/80) and non-immune stromal cells (zsGreen<sup>neg</sup>CD45<sup>neg</sup>), isolated by fluorescence activated cell sorting from established metastatic livers at day 14 after treatment with GEM or untreated (CTR). Bar graph shows relative expression of Gas6 in cells derived from GEMtreated mice and untreated mice (data are from three independent experiments; mean±SEM). (F–H) Representative images (F) of myeloperoxidase (MPO) and Gas6 staining using RNAscope in serial sections from metastatic livers derived from untreated (CTR) or GEM treated mice (n=3 mice/ group). Arrowheads indicate Gas6+ staining in neutrophil-rich areas. Scoring of Gas6 signal per field of view (G) and MPO staining quantification (H). (I–K) Metastatic tumours in livers of the spontaneous mouse pancreatic cancer model Kras<sup>G12D</sup>;Trp53<sup>R172H</sup>;Pdx1-Cre (KPC mice) treated with Gemcitabine (KPC Gem) or left untreated (KPC Ctr) were isolated and analysed (n=3 mice/group). Representative images (I) of MPO and Gas6 staining using RNAscope in serial sections from metastatic tissue sections. Arrowheads indicate Gas6+ staining in neutrophil-rich areas. (J) Scoring of Gas6 signal per field of view and (K) MPO staining quantification. (L, M) Peripheral blood neutrophils were isolated from metastatic PDAC patients during their first cycle of gemcitabine treatment and GAS6 mRNA levels were assessed by real time PCR. Schematic illustration of treatment regimen and patient blood sample collection (L). Quantification of data (M) (BL=baseline, prior treatment) (n=2 patients). Scale bar=50 µM. Data are presented as mean±SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate p values. \*P<0.05; \*\*p<0.01; \*\*\*p<0.001. ANOVA; analysis of variance; H, healthy liver; IHC, immunohistochemistry; M, metastases; n.s., not significant; PDAC, pancreatic ductal adenocarcinoma.

of neutrophils (figure 4B,C). Thus, to test this hypothesis, we isolated metastasis infiltrating neutrophils from metastatic livers from mice treated with saline (control) or gemcitabine, and cocultured them with gemcitabine pretreated pancreatic cancer cells under anchorage independent growth conditions ex vivo. Gemcitabine-treated pancreatic cancer cells were unable to form colonies (figure 4D; online supplemental figure S4K). Strikingly, coculturing of neutrophils isolated from gemcitabine treated metastatic livers with gemcitabine-treated cancer cells enabled the cancer cells to grow and form colonies, while neutrophils isolated from control (saline treated) metastatic lesions were unable to promote cancer cell proliferation (figure 4D). In contrast, while metastases derived macrophages were also able to significantly increase cancer cell colony formation, the macrophage-growth promoting functions were Gas6-independent, unaffected by gemcitabine and markedly less potent compared with neutrophils derived from gemcitabine treated mice (figure 4D; online supplemental figure S4L). In agreement with these findings, in vivo, Ki67<sup>+</sup> cancer cell numbers were reduced in macrophage depleted mice independent of their treatment (online supplemental figure S4M, N). Taken together, these data show that gemcitabine treatment makes neutrophils acquire a promitogenic capacity that promotes cancer cell proliferation.

We next aimed to understand how neutrophils promote cancer cell proliferation. To achieve this goal, we performed RNA sequencing of metastasis infiltrating neutrophils isolated from saline treated metastatic livers (Ly6 $G_{Ctr}$ ) and gemcitabine treated metastatic livers (Ly6G<sub>Gem</sub>) (online supplemental figure S4O). Differently expressed genes were first filtered for GO terms extracellular and receptor ligand activity. Among the resulting n=141 genes, we identified Growth Arrest Specific protein 6 (Gas6) as one of the top upregulated genes in neutrophils derived from gemcitabine treated metastatic lesions compared with control metastatic lesions (online supplemental tables S1, S2). Gas6 and its main receptor AXL are overexpressed in pancreatic cancer and their expression correlates with poor prognosis.<sup>28</sup> <sup>29</sup> Gas6/AXL signalling in cancer cells is associated with tumour cell proliferation, epithelial mesenchymal transition and metastases.<sup>30 31</sup> Subsequent analysis of *Gas6* expression in flow cytometry sorted neutrophils, non-immune stroma cells, macrophages and cancer cells confirmed that within the metastatic tumour microenvironment, neutrophils markedly upregulate Gas6 expression in response to gemcitabine treatment and neutrophils are the main source of Gas6 after gemcitabine withdrawal (figure 4E; online supplemental figure S5A).

In agreement with these findings, we found a marked upregulation of Gas6 expression levels in neutrophil-rich areas proximate to cancer cells after gemcitabine withdrawal in serial tissue sections derived from experimental (figure 4F–H; online supplemental figure S5B, D) and spontaneous hepatic metastatic lesions (figure 4I–K; online supplemental figure S5C, D). In contrast, chemotherapy withdrawal did not increase Gas6 levels in metastatic livers from neutrophil-depleted mice (online supplemental figure S5E, F). Moreover, we only found an increase of Gas6 expressing neutrophils in tumour-bearing livers, but not in tumour-free lung tissues, suggesting Gas6 expressing neutrophils preferentially accumulate at the metastatic tumour site (online supplemental figure S5G, H).

We observed the same changes when we treated metastatic tumour-bearing mice with nab-paclitaxel or FOLFIRINOX, both commonly used chemotherapy regimens in PDAC<sup>25</sup> (online supplemental figure S5I). With all chemotherapeutic treatments, metastatic tumour burden temporarily decreased but was followed by metastatic relapse (online supplemental figure S5J) which was

accompanied by an influx of Gas6-expressing neutrophils into metastatic lesions (online supplemental figure S5K-N). These results suggest that the increased accumulation of Gas6-expressing neutrophils in relapsed metastatic lesions after chemotherapy treatment occurs in response to different chemotherapeutic treatment regimens and is therefore not agent specific. We next assessed Gas6 expression in circulating neutrophils in patients with metastatic pancreatic cancer and in our mouse metastases model. We collected patient blood samples prior (baseline) and after (week 4) their first cycle of gemcitabine treatment (figure 4L). We found that Gas6 expression increased in circulating neutrophils 4 weeks after the first dose of treatment (figure 4M; online supplemental figure S6A). Similarly, in the preclinical mouse model, Gas6 expression was increased in circulating murine neutrophils after gemcitabine withdrawal (online supplemental figure S6B). Since the release of NETs by apoptotic neutrophils has been shown to promote pulmonary metastatic outgrowth in breast cancer models,<sup>16</sup> we also analysed the presence of apoptotic (TUNEL+) neutrophils in liver metastases. However, we could only detect a few apoptotic neutrophils within liver metastases and their numbers remained unaffected by gemcitabine treatment (online supplemental figure S6C, D). To assess the biological importance of Gas6 in promoting regrowth of metastatic cancer cells, we next isolated metastasis infiltrating neutrophils from gemcitabinetreated tumour-bearing mice (Ly6G<sub>Gem</sub>) and cocultured those neutrophils with gemcitabine treated cancer cells in the presence or absence of a Gas6 neutralising antibody. We found that Gas6 secretion from neutrophils promotes cancer cell regrowth, in fact, the addition of a neutralising Gas6 antibody abolished the promitogenic effect of neutrophils (figure 5A). Next, we tested whether Gas6 is sufficient to promote the regrowth of gemcitabine treated cancer cells. We pretreated human Panc1 cells and mouse derived KPC cells with gemcitabine and measured regrowth of the cancer cells in the presence or absence of recombinant Gas6. Addition of recombinant Gas6 was sufficient to promote the regrowth of gemcitabine treated human (figure 5B,C) and mouse pancreatic cancer cells (figure 5D). To further test the role of Gas6 in cancer cell regrowth in metastatic livers after gemcitabine treatment in humans, we generated precision cut liver slices (PCLS) from fresh liver biopsies from treatment naïve metastatic PDAC patients. Next, PCLS were treated ex vivo with gemcitabine for 24 hours, washed, and further cultured in the presence or absence of recombinant Gas6 (figure 5E). PCLS were embedded and we assessed the presence of metastatic cancer cells (Muc1<sup>+</sup>) and proliferating cells (Ki67<sup>+</sup>). In line with our previous colony formation experiments, we found an increase in proliferating metastatic cancer cells (MUC1<sup>+</sup>Ki67<sup>+</sup>) in gemcitabine treated PCLS cultures supplemented with recombinant Gas6 compared with gemcitabine treated PCLS cultures lacking recombinant Gas6 (figure 5F,G; online supplemental figure S6E, F). These experiments suggest that Gas6 is sufficient to promote the regrowth of gemcitabinetreated pancreatic cancer cells not only in PDAC cell lines in vitro, but also in patient-derived metastatic liver samples ex vivo. Taken together, these findings show that (1) the infiltration of neutrophils in metastatic lesions after chemotherapeutic treatment leads to metastatic relapse in vivo, (2) Gas6 expression is highly upregulated in neutrophils after chemotherapy and (3) Gas6 promotes regrowth of gemcitabine-treated pancreatic cancer cells.

# Blockade of Gas6/Axl signalling axis restrains metastatic relapse after gemcitabine treatment

Gas6 is a ligand of the TAM receptor family (Tyro3, AXL, Mer) and binding of Gas6 to a TAM receptor results in its

Gut: first published as 10.1136/gutjnl-2021-325272 on 12 January 2022. Downloaded from http://gut.bmj.com/ on November 11, 2022 at University of Glasgow. Protected by copyright



**Figure 5** Gas6 is necessary for neutrophil-mediated cancer cell regrowth after gemcitabine treatment. (A) Quantification of colony formation assay of gemcitabine treated KPC<sup>luc/zsGreen</sup> cells in the presence or absence of Gas6 neutralising antibody (anti-Gas6) with or without metastasis infiltrating neutrophils (LyGG<sub>Gem</sub>) isolated from mice treated with gemcitabine. Bar graph shows fold upregulation of bioluminescence imaging (BLI) signal compared with gemcitabine-treated KPC<sup>luc/zsGreen</sup> cells alone (three independent experiments; mean±SEM). (B–D) Colony formation assay of gemcitabine-treated human Panc1 and murine KPC<sup>luc/zsGreen</sup> cells in the presence or absence or recombinant Gas6 (rGas6). (B) Representative images of Panc1 colonies. (C) Quantification of colony numbers (fold change compared with untreated Panc1 cells) (three independent experiment; mean±SEM). (D) Quantification of BLI signal from KPC<sup>luc/zsGreen</sup> colonies (fold change compared with untreated KPC cells) (three independent experiment; mean±SEM). (E–G) Schematic illustration of experiment (E): Human precision cut liver slices (hPCLSs) were initially treated with gemcitabine for 24 hours then cultured in the presence or absence of rGas6 for the following 24 hours. hPCLSs were assessed by MUC1 (cancer cell marker) and Ki67 immunofluorescent staining (proliferation marker). (F) Representative if images and (G) quantification of proliferating Ki67 + tumour cell frequency in ex vivo treated hPCLS (n=5 patient biopsies). Arrowheads indicate Ki67 + cancer cells. Scale bar 50 µM. data are presented as mean±SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate p values. \*P<0.05; \*\*p<0.01; \*\*\*p<0.001. ANOVA, analysis of variance; H, healthy liver; M, metastases; n.s., not significant.

phosphorylation and activation.<sup>32</sup> The TAM receptor AXL is highly expressed by pancreatic cancer cells and its activation is associated with poor prognosis.<sup>28</sup> We hypothesised that Gas6 expressing neutrophils activates AXL on disseminated pancreatic cancer cells, thereby enabling cancer cell regrowth. To test this, we assessed AXL phosphorylation in metastatic lesions of mice treated with gemcitabine or saline and in the presence or absence of neutrophils. In agreement with our hypothesis, we found that in metastatic lesions derived from gemcitabine treated mice the level of AXL phosphorylation (pAXL) in disseminated cancer cells was markedly increased compared with control (saline treated) metastatic tumour lesions (figure 6A). Importantly, depletion of neutrophils (aLy6G) abolished AXL activation on disseminated cancer cells (figure 6A,B). These data confirm that AXL receptor activation on disseminated cancer cells requires the presence of neutrophils. We next tested whether pharmacological blockade of the Gas6/AXL signalling pathway using warfarin provides a therapeutic benefit when combined with chemotherapy. Gas6 belongs structurally to the family of plasma vitamin K-dependent proteins and its biological function is dependent on Y-carboxylation, a process that can be blocked

using warfarin<sup>33 34</sup> (figure 6C). Liver metastasis was induced by intrasplenic implantation of KPC cells. After 3 days, mice-bearing micrometastatic lesions were treated with gemcitabine or saline (control) and, 4 days later, we started the treatment with warfarin (figure 6D) to avoid interference with the initial anti-tumour immune response previously observed (figure 2). Gemcitabine or warfarin as monotherapies did not affect hepatic metastatic tumour burden. However, gemcitabine treatment followed by subsequent warfarin administration markedly reduced regrowth of metastatic lesions (figure 6E,F). As expected, warfarin treatment abolished the previously observed increase in AXL activation (pAXL) in cancer cells in the gemcitabine treated cohort (figure 6G,H) and cancer cell proliferation (Ki67<sup>+</sup>) rates were reduced (online supplemental figure S7A, B), while neutrophil numbers remained unchanged (online supplemental figure S7C, D). We and others previously showed that Gas6/AXL signalling also inhibits NK cell activation and warfarin treatment increases NK cell activation and reduces pulmonary metastasis in pancreatic cancer.<sup>35 36</sup> Thus, we assessed the infiltration of NK cells in hepatic metastatic lesions using the NK activation marker NKp46. We found that NK cell numbers in the hepatic



Figure 6 Blockade of the Gas6/Axl signalling pathway via warfarin inhibits metastatic relapse after gencitabine treatment. (A–B) Representative images of pAXL staining in liver tissue sections derived from naïve mice or metastasis bearing mice treated with saline (control) or treated with gemcitabine alone (GEM) or GEM +  $\alpha$ Ly6G (A). Quantification of pAXL + tumour cell frequency (B) (n=3 mice/group). (C) Schematic illustrating Gas6/ Axl blockade via warfarin. (D–H) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc</sup>/<sup>asGreen</sup> cells. At day 3, mice were treated with gemcitabine (GEM) or saline control (saline), at day 7 mice were treated with warfarin (war) or left untreated (CTR). (D) Schematic illustration of the experiment. (E, F) Representative images of BLI signal detected in tumour-bearing livers ex vivo (E) and quantification of tumour burden by ex vivo bioluminescence imaging (BLI) (F) (n=3 mice/group). (G, H) Quantification of pAXL + tumour cell frequency (G) and representative images of pAXL staining of metastatic tumour lesions (H). Arrowheads indicate metastatic cancer cells staining positive for pAXL (n=3 mice/group). (I, J) Primary tumour burden was induced by orthotopic implantation of KPC<sup>luc</sup>/<sup>zsGreen</sup> cells into the pancreas. At day 8, cohorts were treated with GEM or saline control. Treatment with warfarin started at day 12. Livers were resected at day 19 and assessed for metastatic tumour burden (n=5 mice/group). (I) Schematic illustration of the experiment. (J) Quantification of tumour burden by ex vivo BLI (n=5 mice/group). (K–P) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. At day 3, mice were treated with GEM or saline, at day 7 mice were treated with R428 or control vehicle. (K) Schematic illustration of experiment. (L)Change in tumour burden was guantified by ex vivo BLI (n=4 mice/group). (M, N) Quantification of pAXL+ tumour cell frequency (M) and representative images (N). (O, P) Quantification of Ki67+ tumour cell frequency (O) and representative images (P). Arrowheads indicate metastatic cancer cells staining positive for pAXL (N) or Ki67 (P). (Q) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. At day 3, mice were treated with GEM, at day 4, mice were treated with  $\alpha$ Ly6G, at day 7 mice were treated with R428 or control vehicle until end point (day 14). Quantification of tumour burden by ex vivo BLI imaging (n=5 mice/group). (R) Liver metastasis was induced by intrasplenic implantation of 1×10<sup>6</sup> KPC<sup>luc</sup>/<sup>zsGreen</sup> cells. At day 3, animals were treated with GEM or saline control (saline). From day 7, mice were treated with warfarin or left untreated. Survival analysis of gemcitabine, gemcitabine/warfarin, warfarin and saline treated mice-bearing liver metastasis; log-rank (Mantel-Cox) test, p=0.0456 (GEM vs Gem/War). Median survival for saline was 36.5 days (n=6 mice), for warfarin 32.5 days (n=6 mice), gemcitabine 32.5 days (n=6 mice) and gemcitabine/warfarin 42 days (n=6 mice). Scale bar 50 µM. Data are presented as mean±SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate p values. \*P<0.05; \*\*p<0.01; \*\*\*p<0.001. ANOVA; analysis of variance; H, healthy liver; M, metastases; n.s., not significant.

metastatic lesions were very low and not affected by warfarin (online supplemental figure S7E, F).

We next assessed whether warfarin is capable of inhibiting metastatic growth of disseminated cancer cells in a spontaneous metastasis model of pancreatic cancer after chemotherapy treatment. To test this, KPC derived cells were orthotopically implanted into the pancreas. At day 8, after the establishment of primary tumours, animals were treated with gemcitabine, followed by warfarin administration (figure 6I). Gemcitabine and warfarin administration alone, or as combinatorial therapy, did not affect primary tumour burden (online supplemental figure S7G, H). Similar to the experimental metastasis model, gemcitabine and warfarin treatment as monotherapies did not affect metastatic tumour burden compared with control treated mice. However, the presence of warfarin following gemcitabine treatment significantly reduced metastatic tumour burden (figure 6]), AXL activation (pAXL) and proliferation (Ki67<sup>+</sup>) of disseminated cancer cells (online supplemental figure S7I-L), but immune cell infiltration remained unchanged by warfarin (online supplemental figure S7M-Q). To further confirm that Gas6 mediates metastatic relapse through AXL activation and not through one of its other TAM receptors (Tyro3, Mer), we next tested the effect of the AXL inhibitor R428 on metastatic relapse after treatment (figure 6K). In line with our previous findings, pharmacological inhibition of AXL after gemcitabine withdrawal significantly reduced metastatic relapse (figure 6L; online supplemental figure S8A, B), AXL (pAXL) activation and proliferation (Ki67+) of metastatic cancer cells (figure 6M–P), while immune cell infiltration remained unaffected (online supplemental figure S8C-G). Consistent with our findings that metastasis associated neutrophils are a main source of Gas6 expression and are critical for the activation of AXL after chemotherapy withdrawal, AXL inhibition in neutrophil depleted mice did not show any additional therapeutic benefit compared with neutrophil depletion alone (figure 6Q). Moreover, pharmacological inhibition of the Gas6/AXL signalling pathway with warfarin after gemcitabine treatment significantly increased overall survival of mice with liver metastases (figure 6R).

Thus, our findings provide evidence that targeting the AXL receptor is sufficient to reduce metastatic relapse after gemcitabine treatment.

# Chemotherapy treatment upregulates the expression of *Cxcl1* and *2* in pancreatic cancer cells which promotes neutrophil recruitment to the tumour site

Chemotherapies often show adverse side effects in patients, including a transient reduction of immune cell populations caused by the toxicity of the drugs.<sup>37</sup> Hence, we further analysed neutrophil numbers in the peripheral blood of liver metastasis bearing patients and mice in response to chemotherapy treatment. As expected, a single dose of chemotherapeutic agents was sufficient to induce a transient reduction in neutrophil numbers in the blood of patients and mice, while 2 weeks after the last dose of treatment, neutrophil numbers rebounded (online supplemental table S3; online supplemental figure S9A, B). To explore the mechanism promoting the accumulation of neutrophils at the metastatic site in response to chemotherapy after rebound, we next tested whether pancreatic cancer cells treated with chemotherapeutic drugs upregulate the expression of chemokines known to promote neutrophil migration, including CXCL1, 2, 5 and 8 (expressed in human only).<sup>38</sup> We found that in human and mouse pancreatic cancer cells, the chemokines Cxcl1 and Cxcl2 are highly upregulated in response

to gemcitabine treatment (figure 7A). The upregulation of Cxcl1,2 by cancer cells in response to gemcitabine treatment was further confirmed in flow cytometry sorted disseminated pancreatic cancer cells (figure 7B). Interestingly, we also found that within the metastatic site, macrophages are a substantial source of Cxcl1 and 2 expression but their expression levels remained unaffected in response to gemcitabine (figure 7B). Since CXCL1,2,5 and 8 bind to the chemokine receptor CXCR2 which is associated with neutrophil recruitment to tumours,<sup>8</sup> we first confirmed that Cxcr2 is indeed highly expressed in metastasis infiltrating neutrophils (figure 7C) and that CXCR2 expression is not affected by gemcitabine treatment (figure 7D; online supplemental figure S9C). Next, we tested whether pharmacological blockade of CXCR2 was sufficient to ablate neutrophil migration. As expected, while tumour conditioned media generated from gemcitabine-treated human and mouse pancreatic cancer cells significantly increased neutrophil migration in vitro, compared with control conditioned media, the presence of the CXCR2 inhibitor SB225002 inhibited neutrophil migration (figure 7E,F; online supplemental figure S9D, E). Moreover, recombinant CXCL2 was sufficient to induce human and mouse neutrophil migration (online supplemental figure S9F, G). In vivo, pharmacological inhibition of CXCR2 (figure 7G) significantly inhibited metastatic relapse after gemcitabine withdrawal (figure 7H; online supplemental figure S9H) and reduced the accumulation of neutrophils at the metastatic site (figure 7I; online supplemental figure S9I). Taken together, these results show that the neutrophil attracting cytokines Cxcl1 and 2 are highly expressed in metastatic livers in response to gemcitabine withdrawal and this favours CXCR2-dependent recruitment of neutrophils at the hepatic metastatic site.

## Chemotherapy treatment induces accumulation of Gas6 expressing neutrophils in liver metastases of patients with stage IV colorectal cancer

Like pancreatic cancer, colorectal cancer (CRC) frequently metastasises to the liver.<sup>39</sup> While in pancreatic cancer, liver biopsies are only taken prior chemotherapeutic intervention for diagnostic purpose,<sup>25</sup> in CRC, chemotherapy is often the standard-of-care treatment for patients with stage IV CRC (patients with liver metastasis), prior to their metastatic liver surgical resection.<sup>40</sup> These differences in patient care provided us an opportunity to collect liver samples from patients with stage IV CRC treated with chemotherapy, and to analyse whether metastatic liver tumours from patients with CRC show an increase in neutrophil infiltration and Gas6 expression in response to chemotherapy. We analysed resected metastatic liver tumours from untreated patients with stage IV CRC, and from patients with stage IV CRC treated with capecitabine or oxaliplatin. We found that metastatic tumour cells (cytokeratin 19<sup>+</sup>) in both chemotherapytreated patient cohorts were surrounded by higher numbers of neutrophils (MPO<sup>+</sup>) compared with the untreated patient cohort (figure 8A,B). Analysis of serial tissue sections revealed increased levels of Gas6 +expression in neutrophil-rich areas in the treated patients (figure 8C,D; online supplemental figure S9J). To further confirm that neutrophils are indeed a major source of Gas6 expression in human liver metastases after chemotherapy treatment, we enzymatically disaggregated fresh liver samples from chemotherapy-treated patients with CRC into single cell suspensions and isolated neutrophils, macrophages, fibroblasts and cancer cells by flow cytometry-based cell sorting (figure 8E; online supplemental figure S9K). Subsequent gene expression analysis confirmed that neutrophils are the cells



**Figure 7** Chemotherapy treatment upregulates the expression of the neutrophil chemo-attractants CXCL1 and 2 in disseminated tumour cells. (A) Heatmap depicting *Cxcl1, 2, 5, 8* mRNA expression levels assessed by real time PCR in KPC (murine) and PANC-1 (human) pancreatic cancer cell lines untreated (CTR) and gemcitabine treated (GEM) (three independent experiments; mean $\pm$ SEM). (B–D) Liver metastasis was induced by intrasplenic implantation of KPC cells. Cohorts were treated at day three with saline (CTR) or GEM (n=3 mice/group). Cancer cells, macrophages and neutrophils were isolated from metastatic lesions at day 14 by FACS. (B) Quantification of *Cxcl1* and *2* mRNA levels by real time PCR in disseminated KPC cancer cells and macrophages (three independent experiments; mean $\pm$ SEM). (C) Heatmap depicting gene expression analysis of CXCR family receptors (*Cxcr1, 2, 3, 4*) by metastasis infiltrating neutrophils. (D) Quantification of CXCR2 expression levels by flow cytometry on neutrophils isolated from tumour-free livers (naïve) and liver metastases derived from saline (CTR) or gemcitabine (GEM) treated mice (n=3 mice/group). (E, F) Quantification of murine (E) and human (F) neutrophil migration in the presence and absence of CXCR2 inhibitor SB225002 (iCXCR2) towards tumour conditioned media (TCM) generated from pancreatic cancer cells (KPC and Panc1, respectively) exposed to gemcitabine (TCM<sub>Gem</sub>) or control (TCM<sub>ctr</sub>) (three independent experiments; mean $\pm$ SEM). (G–I) Liver metastasis was induced by intrasplenic implantation of KPC cells. Cohorts were treated at day three with GEM or saline (CTR). From day 4 mice were treated with SB225002 (iCXCR2) until endpoint (day 14). (H) Quantification of tumour burden by ex vivo BLI (n=5 mice/group). (I) Flow cytometry quantification of neutrophil frequency in metastatic livers at endpoint. Data are presented as mean $\pm$ SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate p values. \*P<0.05; \*\*p<0.01; \*\*\*p<0.001; ANOVA, analysis of variance; n.s., not

expressing the highest levels of *GAS6* in patient liver metastases after chemotherapy cessation (figure 8F). Taken together, these findings suggest that chemotherapy-induced neutrophil accumulation and upregulation of Gas6 also occurs in liver metastasis of patients with CRC and thus, targeting Gas6 might improve therapeutic interventions in patients with pancreatic cancer and CRC.

## DISCUSSION

Our findings demonstrate that standard cytotoxic chemotherapy temporarily restrains metastatic PDAC progression but, also induces cellular changes in the metastatic tumour microenvironment which subsequently promote metastatic relapse. Specifically, we found that chemotherapy induces the expression of neutrophil chemoattractants in tumour cells and subsequent recruitment and infiltration of Gas6 expressing neutrophils to the liver in a CXCR2-dependent manner. Neutrophil-derived Gas6 then activates the receptor tyrosine kinase AXL on metastatic cancer cells and promotes metastatic growth in the liver (figure 8G). In this study, we also show that pharmacological inhibition of Gas6/AXL signalling in combination with chemotherapy ablates metastatic relapse, thereby providing the rationale for further evaluating this therapeutic strategy for PDAC patients.



**Figure 8** Chemotherapy treatment induces accumulation of Gas6 expressing neutrophils in liver metastases of patients with stage IV colorectal cancer. (A–D) Tissue sections from metastatic livers derived from treatment naïve patients with stage IV colorectal cancer (CRC) (n=5), and patients undergone treatment with oxaliplatin (n=3) or capecitabine (n=4) were stained for cancer cells (CK19), neutrophils (MPO) and *GAS6*. (A, B) Representative images of CK19 and MPO staining of serial sections (A) and quantification of data (B). (C, D) Representative images of *GAS6* and MPO staining of serial sections (C) and quantification of data (D). Arrowheads indicate *GAS6+* staining in neutrophil-rich areas. (E, F) Liver biopsies were collected from metastatic CRC patients post-FOLFOX treatment. Cell populations were isolate by FACS for gene expression analysis. (G) Schematic illustration of experiment and (H) quantification of *GAS6* mRNA levels by real-time PCR in neutrophils, macrophages, fibroblast and cancer cells (n=3 patient samples). (G) Schematic depicting the role of neutrophil-derived Gas6 in hepatic metastatic tumour regrowth after chemotherapy in pancreatic cancer. Chemotherapy induces the expression of the neutrophil-chemoattractants CXCL1 and 2 by disseminated cancer cells. On treatment withdrawal, neutrophils are recruited to the liver and express high levels of Gas6. Neutrophil-derived Gas6 activates AXL receptors on disseminated cancer cells and promotes their regrowth after chemotherapeutic treatments. Depletion of neutrophils or inhibition of Gas6/AXL signalling axis inhibits metastatic regrowth of pancreatic cancer cells. Scale bar 50  $\mu$ M. Data are presented as mean±SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate p values. \*P<0.05; \*\*p<0.01; \*\*\*p<0.001. ANOVA, analysis of variance; H, healthy liver; M, metastases; n.s., not significant.

Primary pancreatic tumours are largely refractory to chemotherapeutic treatments and surgical resection remains the only curative treatment option.<sup>25</sup> Reasons for this include the excessive stromal compartment, hypovascularisation and increased interstitial fluid pressure within PDAC tumours, all acting as a barrier for efficient drug delivery.<sup>41</sup> Patients that have undergone surgery also receive adjuvant chemotherapy as standard of care.<sup>25</sup> In a preclinical mouse model of PDAC, adjuvant gemcitabine treatment, after the resection of the pancreatic tumour, inhibited local recurrence at the primary tumour site, but not at the distant metastatic sites.<sup>42</sup> However, the effect of chemotherapy on metastatic tumours, and micrometastases in the adjuvant setting, remains poorly understood. Our results provide evidence that metastatic PDAC lesions are responsive to chemotherapeutic treatment, particularly during early metastatic development where the stromal compartment is less established.<sup>3</sup> In fact, chemotherapy increases cancer cell death and the release of tumour antigens, a critical step in the generation of an antitumour immune response.<sup>43</sup> In agreement with this, we observe in our PDAC metastatic mouse model that chemotherapy induces an initial proinflammatory response with activation of cytotoxic T cells in the metastatic liver niche. Although this proinflammatory response was only transient, it might provide a window of opportunity for the use of immune checkpoint inhibitors in combination with chemotherapy to further boost this initial T cell activation. Similar to what we observed in the adjuvant setting at the hepatic metastatic site in murine models, in the neoadjuvant setting for PDAC patients, chemotherapy has been shown to restore an anti-tumour immune response at the primary site associated with a decrease in immune suppressive myeloid cells,  $T_{reg}$ , and CD163<sup>+</sup> macrophages.<sup>44–45</sup> Immune suppressive myeloid Ly6G<sup>+</sup> cells suppress CD8<sup>+</sup> T cell functions in primary PDAC tumours.<sup>8 46</sup> Our findings suggest that in liver metastases the presence of immune suppressive M2-like macrophages is key to inhibit the initial chemotherapy-induced antitumour immune response. In contrast, neutrophils are dispensable for the maintenance of an immunosuppressive metastatic tumour microenvironment. This might be due to the fact that macrophages are present in livers in a much higher number compared with neutrophils.<sup>3</sup> We and others have previously shown that metastasis associated macrophages can suppress CD8<sup>+</sup> T cell functions in the liver.<sup>12 13</sup> In agreement with these findings, the use of a CSF-1/CSF-1R inhibitors also reduced metastatic relapse after gemcitabine treatment, confirming that macrophages play a key role in controlling the local immune response in liver metastasis. Since macrophages have a high plasticity and their heterogeneity is diverse in the liver, future treatments should focus on targeting macrophage immune-suppressive functions or on inhibiting macrophage polarisation towards an immunosuppressive phenotype. Neutrophils have been associated with different prometastatic functions, mainly at the early steps of the metastatic cascade or even at the premetastatic niche formation.<sup>8 9 18 47</sup> Here, we have identified a novel prometastatic function of neutrophils in promoting metastatic growth of cancer cells after the initial colonisation steps. Mechanistically, we found that cessation of chemotherapy induces the infiltration of neutrophils and that neutrophils activate AXL receptor on metastatic cancer cells, via secretion of the AXL receptor ligand Gas6, leading to cancer cell growth at the metastatic liver. AXL is overexpressed in pancreatic cancer and is associated with increased metastasis and a poor prognosis.<sup>28</sup> The Gas6/AXL signalling pathway regulates several cancer cell autonomous and non-cancer cell autonomous processes.<sup>48</sup> <sup>49</sup> AXL has been found to induce epithelial-mesenchymal transition in pancreatic cancer cells and thereby promote cancer cell migration, invasion, and metastatic spreading in vivo. Moreover, genetic depletion of AXL in pancreatic cancer cells increased their

sensitivity to chemotherapy.<sup>31 33 50</sup> In our studies, we started AXL inhibition after the initial seeding and colonisation step of the liver and after the exposure to gemcitabine treatment. Hence, our data reveal an additional novel role of AXL signalling in promoting the regrowth of cancer cells after chemotherapy in established distant lesions, thereby expanding the potential use of AXL inhibitors to fight pancreatic cancer. We and others have previously shown that Gas6 is also expressed by macrophages and fibroblasts<sup>51</sup> and that inhibition of Gas6/AXL signalling increases NK cell activation and reduces pulmonary metastasis.<sup>35 36</sup> Interestingly, here, we show that at the hepatic metastatic site, basal Gas6 expression levels are very low, but are markedly increased in the liver-stroma of human and mouse in response to chemotherapeutic treatments, and are mainly expressed by the infiltrated neutrophils. Moreover, we found that hepatic metastatic lesions are poorly infiltrated by NK cells and neither their numbers nor their activation status was altered by warfarin or AXL inhibition, suggesting that the multifunctional Gas6/AXL signalling pathway regulates different processes and cell populations in different organs, thereby contributing in many different ways to PDAC metastasis.

Our studies exemplify two therapeutic options to inhibit metastatic relapse by using warfarin or the AXL inhibitor R428 (Bemcentinib). Both agents are currently tested in patients with pancreatic cancer (NCT03536208) and (NCT03649321), respectively. Hence, our findings further strengthen the rationale for targeting Gas6/AXL signalling in the treatment of metastatic PDAC and, in combination with chemotherapy, to reduce the risk of recurrence in the adjuvant setting by preventing the progression of micrometastatic disease. Further studies will be needed to explore the mechanism by which chemotherapy withdrawal induces Gas6 expression in neutrophils, and to test whether Gas6 levels in circulating neutrophils could be also used as a biomarker for predicting the risk of metastatic recurrence in the adjuvant setting.

Our additional analysis of liver metastases from patients with stage IV CRC further suggests that the observed increase in Gas6-expressing neutrophils in response to chemotherapy might not be restricted to pancreatic cancer metastasis, but may also occur in other cancers that metastasise to the liver. Thus, targeting the identified neutrophil/Gas6/AXL axis might also be of relevance for cancers with a high prevalence to spread to the liver, such as CRC, melanoma, breast and lung.<sup>39</sup> In summary, our findings are important and timely as they could help improve in the near future the design of treatment regimens for patients with cancer with liver metastases.

### MATERIALS AND METHODS

Detailed materials and methods can be found in online supplemental section.

**Acknowledgements** We thank the Liverpool Shared Research Facilities and the CRUK Beatson Institute Biological Science Unit for provision of equipment and technical assistance. We thank Vatshala Chandran for technical support. We also thank the patients and their families, as well as the healthy blood donors who contributed with tissue samples and blood donations to these studies.

**Contributors** GB and CR designed and performed most of the experiments, analysed and interpreted the data, and contributed to the preparation of the manuscript. VQ and YA helped with in vivo experiments, MR and EB helped with in vitro studies. DHP performed bioinformatic analysis. FC helped with the analysis and interpretation of tumour biopsies. DHP, RPJ, DV, PG, CMH and RS provided patient samples. JPM provided KPC mice and KPC-derived tissue samples. AM provided conceptual advice, interpreted data and wrote the manuscript. MCS overall conceived and supervised the project, interpreted data, and wrote the manuscript. All authors critically analysed and approved the manuscript.

**Funding** These studies were supported by grants from Cancer Research UK (A25607, A26978, A26979), Medical Research Council (MR/P018920/1) and Pancreatic Cancer Research Fund for MCS, Wellcome Trust (102521/Z/13/Z) and North West Cancer Research Fund for A.M., Cancer Research UK A17196, A2996 and A25233 for JPM.

Competing interests None declared.

Patient consent for publication Not applicable.

**Ethics approval** This study was approved by Human studies using blood and liver tissue samples were approved by the National Research Ethics (NRES) Service Committee North West-Greater Manchester REC15/NW/0477.

Provenance and peer review Not commissioned; externally peer reviewed.

**Data availability statement** All data relevant to the study are included in the article or uploaded as online supplemental information.

**Supplemental material** This content has been supplied by the author(s). It has not been vetted by BMJ Publishing Group Limited (BMJ) and may not have been peer-reviewed. Any opinions or recommendations discussed are solely those of the author(s) and are not endorsed by BMJ. BMJ disclaims all liability and responsibility arising from any reliance placed on the content. Where the content includes any translated material, BMJ does not warrant the accuracy and reliability of the translations (including but not limited to local regulations, clinical guidelines, terminology, drug names and drug dosages), and is not responsible for any error and/or omissions arising from translation and adaptation or otherwise.

**Open access** This is an open access article distributed in accordance with the Creative Commons Attribution 4.0 Unported (CC BY 4.0) license, which permits others to copy, redistribute, remix, transform and build upon this work for any purpose, provided the original work is properly cited, a link to the licence is given, and indication of whether changes were made. See: https://creativecommons.org/ licenses/by/4.0/.

#### ORCID iD

Michael C Schmid http://orcid.org/0000-0002-3445-0013

#### REFERENCES

- Rhim AD, Mirek ET, Aiello NM, et al. EMT and dissemination precede pancreatic tumor formation. Cell 2012;148:349–61.
- 2 Ryan DP, Hong TS, Bardeesy N. Pancreatic adenocarcinoma. N Engl J Med 2014;371:1039–49.
- 3 Nielsen SR, Quaranta V, Linford A, et al. Macrophage-secreted granulin supports pancreatic cancer metastasis by inducing liver fibrosis. Nat Cell Biol 2016;18:549–60.
- 4 Costa-Silva B, Aiello NM, Ocean AJ, *et al*. Pancreatic cancer exosomes initiate premetastatic niche formation in the liver. *Nat Cell Biol* 2015;17:816–26.
- 5 Aiello NM, Bajor DL, Norgard RJ, et al. Metastatic progression is associated with dynamic changes in the local microenvironment. Nat Commun 2016;7:12819.
- 6 Lee JW, Stone ML, Porrett PM, et al. Hepatocytes direct the formation of a prometastatic niche in the liver. Nature 2019;567:249–52.
- 7 Kaczanowska S, Beury DW, Gopalan V, et al. Genetically engineered myeloid cells rebalance the core immune suppression program in metastasis. *Cell* 2021;184:2033–52.
- 8 Steele CW, Karim SA, Leach JDG, *et al*. Cxcr2 inhibition profoundly suppresses metastases and augments immunotherapy in pancreatic ductal adenocarcinoma. *Cancer Cell* 2016;29:832–45.
- 9 Gordon-Weeks AN, Lim SY, Yuzhalin AE, et al. Neutrophils promote hepatic metastasis growth through fibroblast growth factor 2-dependent angiogenesis in mice. *Hepatology* 2017;65:1920–35.
- 10 Kitamura T, Qian B-Z, Pollard JW. Immune cell promotion of metastasis. Nat Rev Immunol 2015;15:73–86.
- 11 DeNardo DG, Ruffell B. Macrophages as regulators of tumour immunity and immunotherapy. Nat Rev Immunol 2019;19:369–82.
- 12 Yu J, Green MD, Li S, *et al*. Liver metastasis restrains immunotherapy efficacy via macrophage-mediated T cell elimination. *Nat Med* 2021;27:152–64.
- 13 Quaranta V, Rainer C, Nielsen SR, et al. Macrophage-derived granulin drives resistance to immune checkpoint inhibition in metastatic pancreatic cancer. Cancer Res 2018;78:4253–69.
- 14 Jaillon S, Ponzetta A, Di Mitri D, et al. Neutrophil diversity and plasticity in tumour progression and therapy. Nat Rev Cancer 2020;20:485–503.
- 15 Park J, Wysocki RW, Amoozgar Z, et al. Cancer cells induce metastasis-supporting neutrophil extracellular DNA traps. Sci Transl Med 2016;8:361ra138.
- 16 Albrengues J, Shields MA, Ng D, et al. Neutrophil extracellular traps produced during inflammation awaken dormant cancer cells in mice. *Science* 2018;361. doi:10.1126/ science.aao4227. [Epub ahead of print: 28 Sep 2018].
- 17 Itatani Y, Yamamoto T, Zhong C, et al. Suppressing neutrophil-dependent angiogenesis abrogates resistance to anti-VEGF antibody in a genetic model of colorectal cancer. Proc Natl Acad Sci U S A 2020;117:21598–608.

- 18 Wculek SK, Malanchi I. Neutrophils support lung colonization of metastasis-initiating breast cancer cells. *Nature* 2015;528:413–7.
- 19 Coffelt SB, Kersten K, Doornebal CW, et al. II-17-producing gammadelta T cells and neutrophils conspire to promote breast cancer metastasis. Nature 2015;522:345–8.
- 20 Gabrilovich DI, Ostrand-Rosenberg S, Bronte V. Coordinated regulation of myeloid cells by tumours. *Nat Rev Immunol* 2012;12:253–68.
- 21 Garrido-Laguna I, Hidalgo M. Pancreatic cancer: from state-of-the-art treatments to promising novel therapies. *Nat Rev Clin Oncol* 2015;12:319–34.
- 22 Groot VP, Blair AB, Gemenetzis G, *et al*. Recurrence after neoadjuvant therapy and resection of borderline resectable and locally advanced pancreatic cancer. *Eur J Surg Oncol* 2019;45:1674–83.
- 23 Jones RP, Psarelli E-E, Jackson R, et al. Patterns of recurrence after resection of pancreatic ductal adenocarcinoma: a secondary analysis of the ESPAC-4 randomized adjuvant chemotherapy trial. JAMA Surg 2019;154:1038–48.
- 24 Nevala-Plagemann C, Hidalgo M, Garrido-Laguna I. From state-of-the-art treatments to novel therapies for advanced-stage pancreatic cancer. *Nat Rev Clin Oncol* 2020;17:108–23.
- 25 Mizrahi JD, Surana R, Valle JW, et al. Pancreatic cancer. Lancet 2020;395:2008–20.
- 26 Sakamoto H, Attiyeh MA, Gerold JM, et al. The evolutionary origins of recurrent pancreatic cancer. Cancer Discov 2020;10:792–805.
- 27 Chen DS, Mellman I. Oncology meets immunology: the cancer-immunity cycle. Immunity 2013;39:1–10.
- 28 Koorstra J-BM, Karikari CA, Feldmann G, et al. The Axl receptor tyrosine kinase confers an adverse prognostic influence in pancreatic cancer and represents a new therapeutic target. *Cancer Biol Ther* 2009;8:618–26.
- 29 Du W, Brekken RA. Does Axl have potential as a therapeutic target in pancreatic cancer? Expert Opin Ther Targets 2018;22:955–66.
- 30 Wu G, Ma Z, Hu W, et al. Molecular insights of Gas6/TAM in cancer development and therapy. Cell Death Dis 2017;8:e2700.
- 31 Du W, Phinney NZ, Huang H, et al. Axl is a key factor for cell plasticity and promotes metastasis in pancreatic cancer. *Mol Cancer Res* 2021;19:1412–21.
- 32 Graham DK, DeRyckere D, Davies KD, et al. The TAM family: phosphatidylserine sensing receptor tyrosine kinases gone awry in cancer. Nat Rev Cancer 2014;14:769–85.
- 33 Kirane A, Ludwig KF, Sorrelle N, et al. Warfarin blocks Gas6-mediated Axl activation required for pancreatic cancer epithelial plasticity and metastasis. Cancer Res 2015;75:3699–705.
- 34 Geng K, Kumar S, Kimani SG, et al. Requirement of gamma-carboxyglutamic acid modification and phosphatidylserine binding for the activation of TYRO3, Axl, and MERTK receptors by growth arrest-specific 6. Front Immunol 2017;8:1521.
- 35 Ireland L, Luckett T, Schmid MC, et al. Blockade of stromal Gas6 alters cancer cell plasticity, activates NK cells, and inhibits pancreatic cancer metastasis. *Front Immunol* 2020;11:297.
- 36 Paolino M, Choidas A, Wallner S, et al. The E3 ligase Cbl-b and TAM receptors regulate cancer metastasis via natural killer cells. Nature 2014;507:508–12.
- 37 Crawford J, Dale DC, Lyman GH. Chemotherapy-Induced neutropenia: risks, consequences, and new directions for its management. *Cancer* 2004;100:228–37.
- Coffelt SB, Wellenstein MD, de Visser KE. Neutrophils in cancer: neutral no more. *Nat Rev Cancer* 2016;16:431–46.
- 39 Tsilimigras DI, Brodt P, Clavien P-A, et al. Liver metastases. Nat Rev Dis Primers 2021;7:27.
- 40 Dekker E, Tanis PJ, Vleugels JLA, et al. Colorectal cancer. Lancet 2019;394:1467–80.
- 41 Feig C, Gopinathan A, Neesse A, *et al*. The pancreas cancer microenvironment. *Clin Cancer Res* 2012;18:4266–76.
- 42 Gürlevik E, Fleischmann-Mundt B, Brooks J, et al. Administration of gemcitabine after pancreatic tumor resection in mice induces an antitumor immune response mediated by natural killer cells. *Gastroenterology* 2016;151:338–50.
- 43 Mellman I, Coukos G, Dranoff G. Cancer immunotherapy comes of age. Nature 2011;480:480–9.
- 44 Michelakos T, Cai L, Villani V, et al. Tumor microenvironment immune response in pancreatic ductal adenocarcinoma patients treated with neoadjuvant therapy. J Natl Cancer Inst 2021;113:182–91.
- 45 Mota Reyes C, Teller S, Muckenhuber A, *et al*. Neoadjuvant therapy remodels the pancreatic cancer microenvironment via depletion of protumorigenic immune cells. *Clin Cancer Res* 2020;26:220–31.
- 46 Bayne LJ, Beatty GL, Jhala N, et al. Tumor-derived granulocyte-macrophage colonystimulating factor regulates myeloid inflammation and T cell immunity in pancreatic cancer. Cancer Cell 2012;21:822–35.
- 47 Szczerba BM, Castro-Giner F, Vetter M, et al. Neutrophils escort circulating tumour cells to enable cell cycle progression. Nature 2019;566:10.1038/ s41586-019-0915-y:553–7.
- 48 Wu G, Ma Z, Cheng Y, et al. Targeting Gas6/TAM in cancer cells and tumor microenvironment. Mol Cancer 2018;17:20.
- 49 Myers KV, Amend SR, Pienta KJ. Targeting Tyro3, Axl and MerTK (TAM receptors): implications for macrophages in the tumor microenvironment. *Mol Cancer* 2019;18:94.

## Pancreas

- 50 Ludwig KF, Du W, Sorrelle NB, et al. Small-Molecule inhibition of Axl targets tumor immune suppression and enhances chemotherapy in pancreatic cancer. Cancer Res 2018;78:246–55.
- 51 Loges S, Schmidt T, Tjwa M, et al. Malignant cells fuel tumor growth by educating infiltrating leukocytes to produce the mitogen Gas6. *Blood* 2010;115:10.1182/blood-2009-06-228684:2264–73.

# SUPPLEMENTAL MATERIAL

Chemotherapy-induced infiltration of neutrophils promotes pancreatic cancer metastasis via Gas6/AXL signalling axis.

# AUTHORS:

Gaia Bellomo<sup>1#</sup>, Carolyn Rainer<sup>1#</sup>, Valeria Quaranta<sup>1</sup>, Yuliana Astuti<sup>1</sup>, Meirion Raymant<sup>1</sup>, Elzbieta Boyd<sup>1</sup>, Ruth Stafferton<sup>1</sup>, Fiona Campbell<sup>1</sup>, Paula Ghaneh<sup>1</sup>, Christopher Halloran<sup>1</sup>, Dean E Hammond<sup>2</sup>, Jennifer P Morton<sup>3</sup>, Daniel Palmer<sup>1</sup>, Dale Vimalachandran<sup>1</sup>, Robert Jones<sup>1</sup>, Ainhoa Mielgo<sup>1</sup> & Michael C Schmid<sup>1\*</sup>

<sup>1</sup> Department of Molecular and Clinical Cancer Medicine, University of Liverpool, Ashton Street, Liverpool, L69 3GE, UK.

<sup>2</sup> Department of Molecular Physiology and Cell Signalling, University of Liverpool, Ashton Street, Liverpool, L69 3GE, UK.

<sup>3</sup>Cancer Research-UK Beatson Institute and Insitute of Cancer Sciences, University of Glasgow, Switchback Road, Glasgow, G61 1BD

<sup>#</sup>Equally contributing authors

\* to whom correspondence should be addressed mschmid@liverpool.ac.uk

1

# SUPPLEMENTARY FIGURE LEGENDS

# Supplementary Figure S1: Characterisation of gemcitabine induced pancreatic cancer cell death *in vitro* and *in vivo*.

(A) Quantification of caspase 3-7 activities in KPC cells untreated or treated with increasing concentrations of gemcitabine (Gem: 10 nM, 50 nM, 100 nM and 200nM) for 24h and 48h (three independent experiments, mean  $\pm$  SEM). One-way ANOVA with Dunnett's multiple comparison test to 0 nM.

(B-I) Liver metastasis was induced by intrasplenic implantation of 1x10<sup>6</sup> KPC<sup>luc/zsGreen</sup> cells. Starting from day 12, animals were treated with gemcitabine (Gem; 100 mg/kg) or saline (Ctr) every 3<sup>rd</sup> day with 4 doses in total.

(B, C) Representative H&E images (B) and quantification (C) of metastatic lesion area after treatment (day 22).

(D, E) Representative H&E images (D) and quantification (E) of metastatic lesion area at human endpoint (HEP) (n= 5 mice/group/time point).

(F, G) Tumour burden assessed *ex vivo* by bioluminescence imaging (BLI) at humane endpoint (HEP) in spleen and liver (n= 14 mice/group) (F) and representative BLI images (G).

(H, I) Representative IHC images (H) and quantification (I) of apoptotic KPC cells staining positive for cleaved caspase 3 (CC3) at HEP (n= 5 mice /group). Arrowheads indicate apoptotic (CC3<sup>+</sup>) cancer cells in ductal structures.

(J, K) Liver metastasis was induced by intrasplenic implantation of 5x10<sup>5</sup> KPC<sup>luc/zsGreen</sup> cells and animals received one dose of gemcitabine (Gem; 100 mg/kg) or saline (Ctr) at day 3. Representative immunofluorescence images (J) and quantification (K) of apoptotic KPC<sup>luc/zsGreen</sup> cells staining positive for TUNEL at day 14 in saline (Ctr) and Gem treated mice (n= 5 mice /group). Arrowheads indicate apoptotic (TUNEL<sup>+</sup>) cancer cells.

(L) Flow cytometry gating strategy for dendritic cell (DC) populations (CD103<sup>+</sup> and CD11b<sup>+</sup>) and macrophages (Macs). Cells were selected by size using SSC-A vs FSC-A; dead cells were excluded using Sytox blue; live cells were gated for CD45<sup>+</sup> cells and subsequentially gated on F4/80 to separate Macs, and CD11c for total DC; dendritic cells were further separated in the CD103<sup>+</sup> and CD11b<sup>+</sup> subpopulations. Macs, CD103<sup>+</sup> and CD11b<sup>+</sup> DCs were gated on zsGreen to detect uptake of KPC cells.

Scale bars = 100  $\mu$ M (B, D); scale bar 50  $\mu$ M (H, J); M = metastases, H = healthy liver; Data are presented as mean ± SEM. Unpaired t-test or ANOVA was used to calculate P values. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001; n.s., not significant.

# Supplementary Figure S2: Gating strategies used for mass cytometry-based analysis of immune cell infiltrates in pancreatic liver metastases in response to chemotherapy.

(A, B) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (Gem; 100mg/kg) or saline (Ctr) at day 3. Metastatic livers were resected at initial response (day 4) and after treatment withdrawal (day 14). Unsupervised down-sampling was performed with viSNE using equal numbers of manually gated viable CD45<sup>+</sup> cells. (A) Representative viSNE maps showing individual markers highlighted in a dot plot on the axes tSNE1 and tSNE2 for gemcitabine treated samples, with each dot representing a single cell. Resulting cell clusters were used to delineate main immune cell populations. (B) Main immune cell populations as shown in (A) were further analyses for expression of surface molecules associated with an activated phenotype (CD69, CD86, MHC class II) (Ctr d4 n= 4 mice, Gem d4 n= 4 mice; Ctr d14 n= 3 mice; Gem d14 n= 4 mice).

# Supplementary Figure S3: CD4<sup>+</sup> T cell numbers are increased during the initial response to gemcitabine, while CD206+ and YM1+ macrophage numbers are reduced.

(A-L) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (Gem; 100mg/kg) or saline (Ctr) at day 3. Metastatic livers were resected at initial response (day 4) and after treatment withdrawal (day 14) (n= 5 mice/group/time point A-F; n= 3 mice/group/time point G-L).

(A) Flow cytometry gating strategy used to assess T cell subpopulations (CD8<sup>+</sup>, CD4<sup>+</sup> and FoxP<sup>+</sup>  $T_{reg}$  cells) and (B-D) quantification of data.

(E) Flow cytometry gating strategy used for active  $(CD69^+)$   $CD4^+$  T cells and (F) quantification of data (n= 5 mice /group).

(G, H) Flow cytometry gating strategy applied (G) and quantification (H) of CD206<sup>+</sup> cells among macrophages (F4/80<sup>+</sup> cells) in metastatic livers of Gem or saline treated animals during initial response (n=3 mice/group).

(I, J) Representative immunohistochemistry images (I) and quantification (J) of YM-1 positive cells in livers of Gem or saline treated mice during initial response (n=3 mice/group).

(K, L) Representative immunohistochemistry images (K) and quantification (L) of YM-1<sup>+</sup> cells in livers of Gem or saline treated mice after withdrawal (n=3 mice/group).

Scale bar 50  $\mu$ M; M = metastases, H = healthy liver; Arrowheads indicate YM-1<sup>+</sup> cells (macrophages). Data are presented as mean ± SEM. Unpaired t-test was used to calculate P values \**P* < 0.05; \*\*\**P* < 0.001; n.s., not significant.

# Supplemental Figure S4: Macrophage-targeted therapy reduces PDAC liver metastasis independent of gemcitabine treatment, while depletion of neutrophils reduced metastatic tumour burden only after gemcitabine withdrawal.

(A-C, F-H) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (Gem; 100mg/kg) or saline (Ctr) at day 3. At day 4, mice were treated with IgG control or  $\alpha$ Ly6G antibody for 2 weeks. (A) Flow cytometry quantification of neutrophils (CD45<sup>+</sup>CD11b<sup>+</sup>Ly6G<sup>+</sup>Ly6C<sup>low</sup> cells) frequency among immune cells (CD45<sup>+</sup> cells) in peripheral blood of Ctr + IgG treated, Ctr +  $\alpha$ Ly6G antibody treated, Gem + IgG treated, and Gem +  $\alpha$ Ly6G antibody treated mice at endpoint (day18) (n= 3 mice/group). (B-C) Tumour burden quantification by BLI of gemcitabine treated mice (B) receiving isotope control antibody (Gem + IgG) or monoclonal antibody targeting neutrophils (Gem +  $\alpha$ Ly6G) (n= 3 mice /group) and saline treated mice (C) receiving isotope control antibody (Ctr + IgG) or monoclonal antibody targeting neutrophils (Ctr +  $\alpha$ Ly6G) (n= 3 mice /group).

(D-H, M, N) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (Gem; 100mg/kg) or saline (Ctr) at day 3. At day 4, mice were treated with IgG control or  $\alpha$ CSF1 antibody for two weeks. Tumour burden quantification by BLI of gemcitabine treated mice (D) receiving isotope control antibody (Gem + IgG) (n= 3 mice /group) or monoclonal antibody targeting macrophages (Gem +  $\alpha$ CSF-1) (n= 4 mice /group) and saline treated mice (E) receiving isotope control antibody (Ctr + IgG) or monoclonal antibody targeting macrophages (Ctr +  $\alpha$ CSF1) (n= 3 mice /group).

(F-H) At endpoint (d18), liver tumours were surgically resected and analysed by flow cytometry for macrophage and neutrophil infiltration. (F) Flow cytometry gating strategy and (G) quantification of macrophages in response to neutrophil-depletion ( $\alpha$ Ly6G) and macrophage-depletion ( $\alpha$ CSF-1) and (H) neutrophil numbers in response to neutrophils-depletion ( $\alpha$ Ly6G) and macrophage-depletion ( $\alpha$ CSF-1).

(I-J) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine (Gem; 100mg/kg) at day 3. At day 4, mice were treated with  $\alpha$ Ly6G or IgG controls for 2 weeks; at day 7, mice were treated with  $\alpha$ CD8 or

4

IgG controls until endpoint (day 14). Flow cytometry quantification of neutrophil (CD45<sup>+</sup>CD11b<sup>+</sup>Ly6G<sup>+</sup>Ly6C<sup>low</sup> cells) frequency among immune cells (CD45<sup>+</sup> cells) in peripheral blood of Ctr + IgG treated, Ctr +  $\alpha$ Ly6G, Ctr +  $\alpha$ Ly6G and  $\alpha$ CD8 antibody treated (I), and of CD8 T cell (CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> cells) frequency among immune cells (CD45<sup>+</sup> cells) in peripheral blood of Ctr + IgG treated, Ctr +  $\alpha$ Ly6G, Ctr +  $\alpha$ Ly6G and  $\alpha$ CD8 antibody treated (I), and of CD8 T cell (CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> cells) frequency among immune cells (CD45<sup>+</sup> cells) in peripheral blood of Ctr + IgG treated, Ctr +  $\alpha$ Ly6G, Ctr +  $\alpha$ Ly6G and  $\alpha$ CD8 antibody treated (I) = 5 mice /group).

(K) Colony formation assay of KPC cells gemcitabine-treated (Gem) or untreated (Ctr). Bar graph shows fold reduction of BLI signal compared to untreated (Ctr) KPC cells (three independent experiments; mean ± SEM).

(L) Quantification of colony formation assay of gemcitabine treated KPC<sup>luc/zsGreen</sup> cells in the presence or absence of Gas6 neutralizing antibody (anti-Gas6) with or without metastasis infiltrating macrophages (F4/80<sub>Gem</sub>) isolated from mice treated with gemcitabine. Bar graph shows fold upregulation of BLI signal compared to gemcitabine-treated KPC<sup>luc/zsGreen</sup> cells alone (three independent experiments; mean  $\pm$  SEM).

(M, N) Representative IHC images (M) and quantification of proliferating Ki67+ tumour cell frequency in metastatic livers (N).

(O) RNA Sequencing of metastasis infiltrating neutrophils isolated by FACS from tumour bearing livers of mice at day 14 after treatment with Gem or untreated (Ctr). Schematic illustration of experiment.

Scale bar 100  $\mu$ M; M = metastases, H = healthy liver; Data are presented as mean ± SEM. s. Unpaired t-test or ANOVA with Bonferroni was used to calculate P values. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001; n.s., not significant.

# Supplementary Figure S5: Standard of care chemotherapeutic treatment regimens induce the infiltration of Gas6<sup>+</sup> neutrophils into metastatic tumours.

(A) Flow cytometry gating strategy utilised for isolation of KPC<sup>zsGreen</sup> cancer cells, stroma cells, neutrophils and macrophages by FACS.

(B) Representative images of myeloperoxidase (MPO) and *Gas6* staining using RNAscope in metastatic tissue serial sections derived from saline (Ctr) or gemcitabine (Gem) treated mice. Arrowheads indicate *Gas6*<sup>+</sup> staining in neutrophil-rich areas. Micrographs are a higher magnification of same images shown in main Figure 4F.

(C) Spontaneous liver metastases from the autochthonous mouse pancreatic cancer model *Kras*<sup>G12D</sup>;*Trp53*<sup>R172H</sup>;*Pdx1*-Cre (KPC mice) treated with gemcitabine (KPC Gem) or left untreated (KPC Ctr) were isolated and analysed. Representative images of MPO and *Gas6* 

5

staining in metastatic tissue sections. Arrowheads indicate *Gas6*+ staining in neutrophil-rich areas. Micrographs are a higher magnification of the same images as shown in main Figure 4I.

(D) Representative images of *low*, *medium* and *high Gas6* expression levels in metastatic bearing liver tissue sections. Intensity threshold parameters: *low*  $\leq$  0.27, *medium* = 0.57-0.77, *high*  $\geq$  0.78.

(E, F) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. Mice were treated with gemcitabine or saline 3 days post cell implantation. Treatment with  $\alpha$ Ly6G or control IgG started at day 4 (n=4 mice/group). Livers were resected after 14 days. Representative images (E) and quantification (F) of *Gas6* staining using RNAscope in metastatic tissue sections from different cohorts after treatment withdrawal. Arrowheads indicate increased *Gas6*<sup>+</sup> staining in metastatic lesions.

(G) Flow cytometry-based quantification of neutrophils among CD45<sup>+</sup> cells in metastatic livers and tumour-free lungs of gemcitabine treated animals compared to saline (Ctr) treated animals during initial response (n=3 mice/group).

(H) Bar graph shows fold induction of *Gas6* expression in neutrophils isolated from metastatic livers or tumour-free lungs derived from Gem treated mice and untreated mice (n=3 mice/group).

(I-N) Liver metastasis was induced by intrasplenic implantation of 5x10<sup>5</sup> KPC<sup>luc/zsGreen</sup> cells and animals were treated with gemcitabine + nab-paclitaxel (Gem-Nab), FOLFIRINOX (Folfx) or saline (Ctr) at day 5. Metastatic livers were resected after treatment withdrawal (day 15) (n=4 mice/group).

(J) Tumour burden was monitored at day 3, 7 and 15 by BLI *in vivo* imaging. Change in tumour burden in Gem-Nab and Folfx treated groups compared to control group.

(K) Quantification of neutrophil frequency in metastatic livers at endpoint.

(L) Quantification of *Gas6* mRNA levels by real time PCR in intra-metastatic neutrophils (Ly6G<sup>+</sup> cells) isolated by FACS from different cohorts after treatment withdrawal.

(M, N) Representative images (M) and quantification (N) of *Gas6* staining using RNAscope in metastatic tissue sections from different cohorts after treatment withdrawal. Arrowheads indicate increased *Gas6*<sup>+</sup> staining in metastatic lesions.

Scale bar 100  $\mu$ M; M = metastases, H = healthy liver. Data are presented as mean ± SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate P values. \*P < 0.05; \*\*P < 0.01; n.s., not significant.

Supplementary Figure S6: Neutrophils upregulate the expression of *Gas6* after gemcitabine treatment, while apoptotic neutrophil numbers remain the same.

(A) Peripheral neutrophils (CD45<sup>+</sup>CD11b<sup>+</sup>CD14<sup>-</sup>CD15<sup>+</sup>CD16<sup>+</sup>) were isolated from metastatic PDAC patient during their first cycle of gemcitabine treatment using Miltenyi MACS express kit and purity was assessed by flow cytometry (99%).

(B-D) Liver metastasis was induced by intrasplenic implantation of 5x10<sup>5</sup> KPC<sup>luc</sup>/<sup>zsGreen</sup> cells and animals were treated with gemcitabine (Gem) or saline (Ctr) at day 3. Metastatic livers were resected after treatment withdrawal (day 14).

(B) Fold induction of *Gas6* expression in neutrophils isolated from peripheral blood (using Miltenyi kit) of gemcitabine treated mice compared to saline treated (Ctr) mice (n=3 mice/group).

(C, D) Representative immunofluorescent images (C) of apoptotic neutrophils staining positive for TUNEL at day 14 and quantification (D) of the data (n = 5 mice/group; white arrowheads indicate apoptotic neutrophils).

(E, F) Human precision cut liver slices (hPCLSs) were initially treated with gemcitabine for 24 hours then cultured in the presence or absence of rGas6 for the following 24 hours. hPCLSs were assessed by MUC1 (cancer cell marker) and Ki67 staining (proliferation marker) in serial sections. Representative images (E) and quantification (F) of proliferating Ki67<sup>+</sup> tumour cell frequency in *ex vivo* treated hPCLS (n=5 patient biopsies). Arrowheads indicate Ki67<sup>+</sup> cells in cancer cells enriched area.

Scale bar 50  $\mu$ M; M = metastases, H = healthy liver. Data are presented as mean ± SEM. s. Unpaired t-test was used to calculate P values. \**P* < 0.05; \*\**P* < 0.01; n.s., not significant.

# Supplementary Figure S7: Combinatorial therapy using gemcitabine and warfarin inhibits metastatic relapse in experimental and spontaneous metastasis models.

(A-F) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. At day 3, mice were treated with gemcitabine (Gem) or saline control (Ctr), at day 7 warfarin treatment was commenced (n=3 mice/group) until endpoint (day 14).

(A, B) Representative IHC images (A) and quantification (B) of Ki67<sup>+</sup> cancer cells in liver tumours at endpoint.

(C, D) Representative IHC images (C) and quantification (D) of MPO<sup>+</sup> neutrophils in liver tumours at endpoint.

(E, F) Representative IHC images (E) and quantification (F) of NKp46<sup>+</sup> NK cells in liver tumours at endpoint. Arrowheads indicate NKp46<sup>+</sup> NK cells.

(G-Q) Primary tumour was induced by orthotopic implantation of 2x10^5 KPC<sup>luc/zsGreen</sup> cells in the pancreas. At day 8, mice were treated with gemcitabine (Gem) or control saline (saline), at day 12 mice were treated with warfarin until endpoint (day19) (n=5 mice/group).

(G, H) Representative images of primary tumour (G) and quantification (H) of tumour mass at endpoint.

(I, J) Representative IHC images (I) and quantification (J) of pAXL<sup>+</sup> cancer cells in liver tumours at endpoint. Arrowheads indicate cancer cells staining positive for pAXL.

(K, L) Representative IHC images (K) and quantification (L) of proliferating Ki67<sup>+</sup> cancer cells in liver tumours at endpoint.

(M-Q) Quantification of neutrophil infiltration (M), macrophages (N, O) and NK cell (P, Q) activation at the metastatic site by flow cytometry.

Scale bar 50  $\mu$ M; M = metastases, H = healthy liver. Data are presented as mean ± SEM. s. ANOVA with Bonferroni was used to calculate P values. \*P < 0.05; \*\*P < 0.01; n.s., not significant.

# Supplementary Figure S8: AXL signalling is necessary for metastatic relapse after gemcitabine treatment.

(A-G) Liver metastasis was induced by intrasplenic implantation of KPC<sup>luc/zsGreen</sup> cells. At day 3, mice were treated with gemcitabine (Gem) or saline (Ctr), at day 7 the treatment with R428 (AXL-inhibitor) was commenced (n=5 mice/group).

(A, B) Representative images (A) of H&E staining and quantification (B) of metastatic area in livers.

(C-E) Quantification of neutrophil infiltration (C) and macrophage activation (D, E) at the metastatic site by flow cytometry.

(F, G) Quantification of NK cell infiltration (F) and NK cell activation (G) at the metastatic site by flow cytometry.

Scale bar 50  $\mu$ M; M = metastases, H = healthy liver. Data are presented as mean ± SEM. Unpaired t-test or ANOVA with Bonferroni was used to calculate P values. \*P < 0.05; \*\*P < 0.01; n.s., not significant.

Supplementary Figure S9: Flow cytometry-based characterisation of primary neutrophils in human and mouse.

(A) Peripheral blood was collected from metastatic PDAC patients during their first cycle of gemcitabine treatment and neutrophils absolute count [per ml] was assessed. (BL= baseline, w1 = one week after treatment, w4= four weeks after treatment) (n=2 patients).

(B) Liver metastasis was induced by intrasplenic implantation of 5x10<sup>5</sup> KPC<sup>luc</sup>/<sup>zsGreen</sup> cells and animals were treated with gemcitabine (Gem), gemcitabine + nab-paclitaxel (Gem-Nab), FOLFIRINOX (Folfx) or saline control (Ctr) at day 5. Peripheral blood was analysed by flow cytometry to assess neutrophil numbers (n=3 mice/group).

(C) Flow cytometry gating strategy to measure CXCR2 expression on murine neutrophils in metastatic liver.

(D, E) Flow cytometry analysis to assess purity of primary mouse and human neutrophils used in migration assay. (D) Purity of neutrophils isolated from murine spleen and (E) human peripheral blood.

(F) Quantification of murine neutrophil migration towards recombinant CXCL2 (rCXCL2) using migration assay (three independent experiments; mean ± SEM).

(G) Quantification of human neutrophil migration towards recombinant CXCL2 (rCXCL2) using migration assay (three independent experiments; mean ± SEM).

(H, I) Liver metastasis was induced by intrasplenic implantation of KPC cells. From day 4, mice were treated with SB225002 (iCXCR2) or saline until endpoint (day14). (H) Quantification of tumour burden by *ex vivo* BLI (n = 5 mice/group). (I) Flow cytometry quantification of neutrophil frequency in metastatic livers at endpoint.

(J) Serial sections from metastatic livers derived from treatment naïve stage IV CRC patients (n=5), and patients undergone treatment with oxaliplatin (n=3) or capecitabine (n=4) were stained for neutrophils (MPO) and *GAS6*. Micrographs are a lower magnification of the same images as shown in main Figure 8C.

(K) Flow cytometry gating strategy used for sorting immune cell populations (neutrophils, fibroblast, macrophages and cancer cells) from CRC patient liver biopsies.

Scale bar 100  $\mu$ M, Data are presented as mean ± SEM. s. Unpaired t-test or ANOVA with Bonferroni was used to calculate P values. \**P* < 0.05; \*\**P* < 0.01; n.s., not significant.

# SUPPLEMENTAR MATERIALS AND METHODS

#### Cells

The murine pancreatic cancer cells KPC FC1199 and KPC FC1245 were kindly provided by the Tuveson laboratory (Cold Spring Harbor Laboratory). KPC cells initially were isolated from PDAC tumour tissues of Kras<sup>G12D/+</sup>; p53<sup>R172H</sup>; Pdx1-Cre mice of a pure C57BL/6 background and authenticated as previously reported [1]. KPC<sup>luc/zsGreen</sup> cells were then generated by using pHIV Luc-zsGreen (gift from B. Welm, University of Utah, Salt Lake City, UT; Addgene plasmid no.39196) lentiviral particle infection. Infected cells were selected for high zsGreen expression levels using flow cytometry cell sorter (ARIA, BD). Human pancreatic cancer cell line PANC1 were purchased from ATCC. All cells were cultured in DMEM+10% FBS and supplemented with 2.5  $\mu$ g/mL Amphotericin B (Sigma). All cells were routinely tested negative for the presence of *Mycoplasma* contamination. None of the cell lines used in this article are listed in the International Cell Line Authentication Committee and NCBI Biosample database of misidentified cell lines.

#### Mice

Six- to eight-week-old female C57BL/6 mice were purchased from Charles River Laboratories. All animal experiments were performed in accordance with current UK legislation under an approved project license PPL P16F36770 (M.C. Schmid). Mice were housed under specific pathogen-free conditions at the Biomedical Science Unit at the University of Liverpool. For tumour studies, female animals aged 6 to 8 weeks were used.

#### Metastasis studies

In the survival studies, liver experimental metastases were induced by implanting 1 x 10<sup>6</sup> KPC<sup>luc/zsGreen</sup> FC1199 in 25 µL of PBS into the spleen of immunocompetent syngeneic C57BL/6 mice using a Hamilton 29 G syringe, as previously described [2, 3]. Tumour bearing mice were treated with gemcitabine-HCI (Selleckchem, 100 mg/kg) starting from day 12 and administrated via i.p. injection in a Q3Dx4 schedule until the humane endpoint, or tumour bearing mice were treated with a single injection at day 3. For selected survival studies macrophages or neutrophils were depleted using colony-stimulating factor 1 receptor (CSF-1R) –neutralizing antibody (BioXCell, clone AFS98) or Ly6G-neutralizing antibody (BioXCell, clone 1A8), respectively. Anti-CSF1R was administered via i.p. injection with an initial dose of 400 µg and subsequent doses every other day containing 200 µg. Anti-Ly6G was administered via i.p. injection with 200 µg every other day. Rat IgG1 (BioXCell; clone

HRPN) was used as isotype control in anti-CSF1 study and Rat IgG2 (BioXCell; clone 2A3) was used as isotype control in the neutrophil depletion studies and in the anti-CSF1R study, respectively. In one survival study tumour bearing untreated and gemcitabine-treated mice were randomized to receive warfarin dissolved at a concentration of 0.5 mg/L in drinking water, from day 7 onwards. The water bottles containing warfarin were replenished every other day.

Liver experimental micro-metastases were induced by implanting 5 x 10<sup>5</sup> KPC<sup>luc/zsGreen</sup> FC1199 in 25 µL of PBS into the spleen of immunocompetent isogenic C57BL/6 mice using a Hamilton 29 G syringe. For selected experiments, macrophages, neutrophils or CD8+ T cells were depleted using colony-stimulating factor 1 (CSF-1) -neutralizing antibody (BioXCell, clone 5A1), Ly6G-neutralizing antibody (BioXCell, clone 1A8) or CD8-neutralizing antibody (clone 2.43) respectively; in one experiment anti-Ly6G and anti-CD8 were used in combination. Anti-CSF1 was administered via i.p. injection with an initial dose of 400 µg and subsequent doses 6 days a part containing 200 µg. Anti-Ly6G was administered via i.p. injection with 200 µg every other day. Anti-CD8 was administered via i.p injection with 200 µg every other day. Rat IgG1 (BioXCell; clone HRPN) was used as isotype control in the anti-CSF1 macrophage depletion studies, Rat IgG2 (BioXCell; clone 2A3) was used as isotype control in anti-Ly6G neutrophil depletion studies and Rat IgG2b (BioXclone LTF-2) in the anti-CD8T cell depletion studies, respectively. Tumour bearing mice were treated with gemcitabine-HCI (100 mg/kg) 3 days post cancer cell implantation via i.p. injection. In one experiment, mice were treated with FOLFIRINOX (Oxaliplatin 2.5 mg/kg, Fluorauracil 25 mg/kg, Irinotecan 25 mg/kg, Folinic acid 50mg/ml) via i.v. or gemcitabine-Nab Paclitaxel (gemcitabine 50 mg/kg via i.p., Abraxane 50 mg/kg via i.v), 5 days post cancer cell implantation. Oxaliplatin, Fluorauracil, Irinotecan were kindly provided by the Liverpool Clatterbridge Cancer Centre. In one experiment gemcitabine-treated mice were dosed with SB225002 (iCXCR2, CXCR2 inhibitor, Sigma, SML0716) starting from day7 via i.p. at a dose of 10 mg/kg daily. In one experiment, untreated and gemcitabine-treated mice were randomized to receive warfarin dissolved at a concentration of 0.5 mg/L in drinking water, from day 7 onwards. The water bottles containing warfarin were replenished every other day. In multiple experiments, untreated and gemcitabine-treated mice and neutrophils-depleted gemcitabine- treated mice were treated with Bemcentinib (R248) (MCE, HY-15150), starting from day 7 post cancer cell implantation until the endpoint. Mice were administered twice per day with 7  $\mu$ g of R248, dissolved in 5% DMSO 95% corn oil, via oral gavage.

# Syngeneic orthotopic pancreatic cancer model

Primary tumour formation was induced by implanting  $2 \times 10^5$  KPC<sup>luc/zsGreen</sup> FC1245 in 30 µL of Matrigel (VWR, 734-0269) into the pancreas of immune-competent syngeneic C57Bl/6 6-to 8-week-old female mice using a Hamilton 29 G syringe. Tumour bearing mice were treated with gemcitabine-HCl (100 mg/kg) 8 days post cancer cells implantation via i.p. injection. Untreated and gemcitabine-treated mice were randomized to receive Warfarin dissolved at a concentration of 0.5 mg/L in drinking water from day 12 until endpoint. The water bottles containing warfarin were replenished every other day.

# Autochthonous model of pancreatic cancer KPC

KPC mice on a mixed background were bred in-house at the CRUK Beatson Institute and maintained in conventional caging with environmental enrichment, access to standard chow and water *ad libitum*. All animal experiments were performed under a UK Home Office licence and approved by the University of Glasgow Animal Welfare and Ethical Review Board. Genotyping was performed by Transnetyx (Cordoba, TN, USA). KPC mice of both sexes were treated with gemcitabine-HCl in saline (100 mg/kg). Mice were monitored 3 times weekly and palpable tumour bearing mice were recruited onto study. Vehicle and gemcitabine were administered by intraperitoneal injection every 3 days. Mice were checked 3 times per week and culled and tissues harvested at humane endpoint.

# Metastatic tumour burden quantification

Liver metastasis and spleen tumour burden was quantified, *in vivo* and *ex vivo*, by assessing bioluminescence signal (BLI; IVIS, Perkin Elmer) generated by KPC<sup>luc/zsGreen</sup> cells. For *in vivo* detection, the BLI signal was measured post i.p. injection of Beetle luciferin (Promega; 3 mg/mouse) and quantified as total flux (photons/sec). Metastases-bearing mice were subject to *in vivo* BLI prior treatment initiation to quantify the change in tumour burden. For *ex vivo* investigation, livers and spleens were surgically removed, covered with luciferin and bioluminescence signal measured. Total flux were quantified using Living Image software v4.5 (Perkin Elmer). In some experiments, mice were euthanized at indicated time points and metastatic tumour burden was assessed by quantifying the size of metastatic lesions in haematoxylin and eosin-stained paraffin-embedded liver sections. Zeiss Axio Observer Light Apotome.2 Microscope and Zeiss ZEN imaging software were used.

# Human studies

Human studies using blood and liver tissue samples were approved by the National Research Ethics (NRES) Service Committee North West – Greater Manchester REC15/NW/0477.

Human blood samples were obtained from healthy donors or from advanced PDAC patients with liver metastasis (all pathologically confirmed) undergoing their first cycle of gemcitabine treatment (one treatment/week during 3 weeks (week 0, 1, 2), followed by two weeks rest (week 4). Blood samples were collected at the following two time points: prior to treatment (baseline, week 0) and week 4. Blood was collected in EDTA coated tubes and immediately processed for neutrophils isolation. Haematology blood results were provided by the Liverpool Clatterbridge Cancer Centre for the following three time points: baseline, week 1 and week4.

Liver biopsies were collected from treatment-naïve advanced PDAC patients with liver metastasis (all pathologically confirmed). Liver biopsies from patients undergoing surgical resection of colorectal liver metastases (treatment naïve and following neoadjuvant chemotherapy with oxaliplatin, FOLFOX, and/or capecitabine) were analysed following appropriate ethical approval (REC 15/NW/0477). All individuals provided informed consent for blood or tissue donations on approved institutional protocols.

# In vitro chemotherapeutic treatment

KPCs and Panc1 cells were seeded in petri dishes (d = 10 cm). At 70-80% confluence, murine and human cells were washed in PBS and treated with 100 nM or 200 nM gemcitabine-HCl, respectively, in DMEM + 2% FBS, for 24 hours.

# CellTiter-Glo® luminescent cell viability assay

KPC cells were seeded as previously described and treated with 10, 20, 50, 100 and 200 nM of gemcitabine-HCl for 24 or 48 hours. Cell viability was assessed by CellTiter-Glo® assay according to manufactures (Promega, G9241).

# Preparation of tumour conditioned media

For generating tumour conditioned media (TCM) from gemcitabine-treated cells (TCM-gem), cells were cultured to reach 70-80% confluency. Murine and human cell lines were washed 3 times with PBS before addition of 100 nM and 200 nM gemcitabine, respectively in DMEM containing 2% FBS for 24 hours. After gemcitabine treatment, cells were washed 3 times with PBS and incubated with serum free media for 24 hours. The supernatant was harvested and filtered to remove debris and dead cells using a 0.22  $\mu$ m syringe filter (Fisher Scientific). The control TCM (TCM-Ctr) was generated by seeding cancer cells and allowing them to grow to 70-80% confluency. KPC FC1199 and Panc1 were incubated with 2 % FBS DMEM for 24 hours; cells were washed and cultured with serum free DMEM for 24 hours. The supernatant was harvested and filtered to remove debris and cultured with serum free DMEM for 24 hours. The supernatant was harvested and filtered to remove debris and cultured with serum free DMEM for 24 hours. The supernatant was harvested and filtered to remove debris and cultured with serum free DMEM for 24 hours. The supernatant was harvested and filtered to remove debris and dead cells using a 0.22  $\mu$ m syringe filter (Fisher Scientific).

## Nanostring technology

Pieces of <3 mm<sup>3</sup> of metastatic liver were submerged in 10 volumes of RNAlater (Qiagen). Extraction of total RNA was performed using the RNeasy Mini kit (Qiagen), following the manufacturer's instructions. The RNA concentrations of samples were measured using a NanoDrop 2000 Spectralphotometer (Thermo Fisher); all RNA samples used in this study exhibited optical density (OD) 260/280 ratios between 1.9 and 2.1 and OD 260/230 ratios above 1.8. A total of 770 immune-related mouse genes were analysed using the nCounter Mouse PanCancer Immune Profiling Panel (NanoString). Data analyses were performed using the manufacturer specific software nSolver™ version 4.0 with the nCounter Advanced Analysis plugin version 2.0. To analyse the activity of regulated pathways, we used pathway scoring with default settings as provided by the software. Pathway scores calculated by the software plugin are based on a sample's gene expression profile using the first principal component analysis of a gene set data, with a positive score corresponding to increased expression of more than half its genes.

#### **RNA** hybridization

RNA *in situ* hybridization (RNAScope, ACD Bio-Techne) was performed on formalin-fixed paraffin-embedded sections using the RNAScope 2.5 HD Detection Kit RED according to manufacturer's instructions. The probe used was directed against mouse Gas6 (cat no 473831) or human Gas6 (cat no 427811). The images, taken using Zeiss Axio Observer Light Apotome.2 Microscope, were analysed trough the software QuPath for automated analysis by assigning a staining score for Gas6 of *low, medium* or *high* to each cell

according to the staining intensity (intensity threshold parameters:  $low \le 0.27$ , medium = 0.57-0.77,  $high \ge 0.78$ ).

## Flow cytometry and cell sorting

Single-cell suspensions from murine metastatic liver tumours were prepared by first selecting macroscopically tumour-rich areas in the liver followed by mechanical and enzymatic disruption in Hanks Balanced Salt Solution (HBSS) with 1 mg/mL Collagenase P (Roche) as previously described [2, 4]. In some occasions, tumour free lungs were resected and processed using the same methods. Briefly, cell suspensions were centrifuged for 5 minutes at 1,200 rpm, resuspended in HBSS, and filtered through a 500 µm polypropylene mesh (Spectrum Laboratories). Cell suspensions were resuspended in 1mL 0.05% trypsin and incubated at 37 °C, for 5 minutes. Cells were filtered through a 70 mm cell strainer (Miltenyi) and resuspended in 0.5% BSA/PBS. Peripheral blood was obtained by performing tail nick and collected in EDTA coated tubes (Sarstedt). Red blood cells were lysed using 1x Red blood cell lysis buffer (Biolegend). Cells were resuspend in 2% BSA/PBS, blocked for 10 minutes on ice with Fc block (BD Pharmingen; clone 2.4 G2) and stained with Sytox-blue viability marker (Life Technologies) and conjugated antibodies against CD45 (clone 30F-11), F4/80 (clone BM8), CD206 (clone C068C2), MHCII (clone M5/114.15.2), CD86 (clone GL-1), CD3ε (clone 145-2C11), CD8α (clone 53-6.7), CD4 (clone RM4-5), NK-1.1 (clone PK136), CD69 (clone H1.2F3), CD11b (clone M1/70), Ly6C (clone HK 1.4), Ly6G (clone 1A8), CXCR2 (clone SA044G4), CD103 (clone 2E7), CD11c (clone 418) and FoxP3 (clone Mf-14) (all purchased from Biolegend). Intracellular staining of FoxP3 staining was performed after surface staining; cells were fixed and permeabilised using FOXP3 Fix/Perm Buffer Set (Biolegend) according to manufacture instructions. Human liver biopsies were collected in cold PBS tubes and processed as previously described for obtaining single cell suspensions. Cells were resuspended in 2% BSA/PBS, blocked using human TrueStain FcX (Biolegend) and then stained with Sytox-blue viability marker (Life Technologies) and conjugated antibodies against CD45 (clone HI30), CD11b (clone M1/70), CD14 (clone HCD14), CD15 (clone HI98), CD16 (clone 3G8) epCAM (clone 9C4), HLA-DR (clone L243) and PDGFRbeta (clone 18A2).

Flow cytometry was performed on a FACS Canto II (BD Biosciences), and fluorescenceactivated cell sorting (FACS) was carried out using FACS Aria IIIu (BD Biosciences). Cells were sorted directly in RLT buffer +  $\beta$ -mercaptoethanol according to the manufacturer's instruction for RNA isolation (Qiagen), or in MACS buffer or DMEM supplemented with 20% FBS for *in vitro* assays.

#### Analysis and quantification of immune cells in liver metastasis by mass cytometry

Single-cell suspensions from murine metastatic liver tumours were prepared as described above. Suspensions were further enriched for immune cells by density gradient centrifugation using Histopaque-1083 (Sigma Aldrich) at 400x g for 30 minutes at room temperature without brakes. The cloudy band/interface containing the cells plus the bottom layer was transferred into new tube and gently washed with PBS. After one wash with PBS, the cell suspension was washed in double-deionised water (ddH2O;  $\geq$  18 $\Omega$ )/Maxpar cell staining buffer (1:2 dilution). The pellet was resuspended in 1 ml of Maxpar cell staining buffer and cells were stained with Cell-ID 195-Cisplatin (Fluidigm) viability marker in Maxpar PBS (Fluidigm) for 5 min. Cells were washed with Maxpar cell staining buffer, blocked with Fc Block (BD Pharmingen, Clone 2.4G2) on ice for 10 min and metal-conjugated antibody cocktail added and incubated for 40 min at 4°C. The antibody cocktail contained <sup>141</sup>Pr-Ly6G (clone 1A8), <sup>142</sup>Nd-CD11c (clone N418), <sup>143</sup>ND-CD69 (clone H1.2F3), <sup>146</sup>Nd-F4/80 (clone BM8), <sup>147</sup>Sm-CD45 (clone 30-F11), <sup>148</sup>Nd-CD11b (clone M1/70), <sup>149</sup>Sm-CD19 (clone 6D5), <sup>152</sup>Sm-CD3ε (clone 145-2C11), <sup>162</sup>Dy-Ly6C (clone HK1.4), <sup>168</sup>Er-CD8α (clone 53-6.7), <sup>169</sup>Tm-TCRb (clone H57-597), <sup>170</sup>Er-NK1.1 (clone PK136), <sup>172</sup>Yb-CD86 (clone GL1), <sup>174</sup>Yb-MHC II (clone M5/114.15.2) and <sup>176</sup>Yb-B220 (clone RA3-6B2). Antibodies were used at the concentrations recommended by the manufacturer. Cells were then washed twice in cell staining buffer and stained with 125 µM Intercalator-191 Ir (Fluidigm) diluted 1:2,000 in Maxpar fix and perm buffer (Fluidigm) overnight at 4°C. Cells were then washed twice in Maxpar cell staining buffer followed by two washes in  $18\Omega$  distilled water (Fluidigm) and resuspended in 0.1X EQ<sup>™</sup> Four Element Calibration Beads (Fluidigm) prior acquisition on the Helios CyTOF system (Fluidigm). Samples were acquired at a rate of <500 events/sec. All generated FCS files were normalized and EQ beads standard [5]. Data analysis was performed using Cytobank software (mrc.cytobank.org, v6.3 and v7.0, Beckman Coulter); manual gating was used to remove debris, identify single cells (191 lr+) and to distinguish between dead cells (195Pt+). viSNE analysis was performed on the data utilizing t-stochastic neighbour embedding (t-SNE) mapping based on high dimensional relationships. Viable CD45+ singlets selected by manual gating were used for viSNE unsupervised clustering using equal sampling. Manual gating was then performed on the viSNE map created to determine cell population percentages.

#### **RNA Sequencing**

Murine neutrophils were isolated from metastasis bearing livers by FACS as described in the previous section. Cells were lysed in RNAlater buffer (Qiagen) and total RNA purification was performed with the RNeasy Kit (Qiagen). Total RNA integrity was assessed using Qubit and Bioanalyzer from Agilent. RNA-Seq libraries were prepared using the NEBNExt polyA selection and Ultra Directional RNA library preparation kit following manufactures. Libraries were PCR amplified and sequenced on an Illumina HiSeq 400 System, generating 150 bp reads. Reads were aligned and gene level counts generated using STAR 2.7.2b using mouse genome 'GRCm38' downloaded from Illumina igenomes [6]. Quality control metrics were checked using FastQC [7]. Genes with counts < 10 were removed and the data was normalised for differences in sequencing depth using the robust fragments per million (FPM) method available in the R package DESeq2[8]. Genes were annotated using KEGG pathways or Gene Ontology terms using the relevant annotation packages available within BioConductor. Normalised fragments per million mapped fragments (FPM) count data were obtained for a total of 12,651 genes. Those with a mean FPM of <1 were filtered out, as were genes with counts for only one condition (control [CTR] or Gem-treated [CTX]), creating a dataset with 9,616 differentially-expressed genes. Genes annotated with the Gene Ontology (GO) Cellular Component terms "cell surface" (GO:0009986) and/or "extracellular region" (GO:0005576) and GO Molecular Function (GOMF) term "receptor ligand activity" (GO:0048018) were extracted for further analysis, to generate supplementary tables 1 and 2.

#### Magnetic bead isolation of human and murine neutrophil and murine macrophages

Murine neutrophils were isolated from liver and spleen, macrophages were isolated from liver. Liver samples were prepared as described above for preparation of flow cytometry samples and Ly6G<sup>+</sup> neutrophils and F4/80<sup>+</sup> macrophages were isolated by positive selection, according to the manufacturer's instructions (Miltenyi). Dissected spleens were dissociated in MAC buffer and passed through a 70  $\mu$ m cell strainer to obtain a single-cell suspension. Cells were centrifugated (400 x *g*), and red blood cells were lysed using 1X Red blood cell Lysis buffer (Biolegend). Single cell suspensions were stained, and Ly6G<sup>+</sup> cells were isolated by positive selection, according to the manufacturer's instructions (Miltenyi). Where human neutrophils were isolated from whole blood, blood was collected in EDTA coated tubes and immediately processed for neutrophil isolation using MACSxpress® Whole Blood Neutrophil Isolation Kit (Miltenyi) according to manufacturer's protocol.

# In vitro colony formation assay

KPC<sup>luc/zsGreen</sup> and parental KPC FC1199 and Panc1 cells were seeded and treated with gemcitabine as described in the previous paragraph. Tumour cells were then washed in PBS and serum starved in DMEM, for 24 hours, except for KPC<sup>luc/zsGreen</sup> which were incubated in 2%FBS/DMEM and 100nM gemcitabine for 24 hours. A 24-well or 48-well plate was coated with a layer of 0.5% agar solution made of phenol-free DMEM without FBS. Tumour cells were strained through a 70µm cell filter to ensure a single cell suspension. The cells were then embedded at a concentration of 10000 or 40000 cells/well in a 0.3% agar mix consisting of phenol-free DMEM supplemented with 2% FBS. A layer of 2% FBS phenol-free DMEM was added on top of the agar matrix. In one experiment, tumour cells were incubated with neutrophils (Ly6G<sup>+</sup> cells) or macrophages (F4/80<sup>+</sup> cells), at a ratio 1:1. Liver metastasis derived neutrophils or macrophages were MACS-sorted (as described in previous section from untreated (Ly6G<sub>CTR</sub> or F4/80<sub>CTR</sub>) or gemcitabine treated (Ly6G<sub>CTX</sub> or F4/80<sub>CTX</sub>) tumour bearing mice and co-cultured with cancer cells for 6 days. In another experiment, the system was supplemented with neutralising anti-gas6 antibody (R&D Systems, AB885). Colony quantification was performed after 5 days, by measuring bioluminescence signal (IVIS, Perkin Elmer) following the addition of 10 µL of Beetle luciferin stock (1:100, final concentration 150µg/mL).

In some experiments, KPC parental FC1199 and Panc1cells were embedded with the addition of 400 ng/ml rGas6 (R&D Systems, 885-GSB-050); rGas6 was added to DMEM top layer and replaced every other day. The colonies were fixed and stained with 0.05% crystal violet in methanol/PBS and the number of colonies per well was counted at day 14 after cell seeding by bright-field microscopy.

## Transwell migration assay

1 x 10<sup>6</sup> human or murine neutrophils were added in serum free DMEM in Transwell inserts with 5  $\mu$ m pore size (Corning) and allowed to migrate to the bottom chamber of the 24-well plate for 15 hours. In one experiment, the bottom chamber was loaded with serum free DMEM +/- rCXCL2 (Peprotech, 250-20A) at a concentration of 50 ng/ml. In another experiment, the bottom chamber was loaded with tumour conditioned media generated from untreated (TCM-ctr) or gemcitabine-treated (TCM-gem) murine or human cancer cell lines (KPC or Panc1) in the presence or absence of SB225002 (iCXCR2, CXCR2 inhibitor, Sigma, SML0716) at a concentration of 25  $\mu$ g/ml. Migrated neutrophils were recovered from the lower chamber and counted by using a haemocytometer.

#### Precision cut liver slicing and ex vivo culture

Precision cut liver slicing was performed as previously reported [9, 10]. Briefly, PDAC patient liver biopsies were embedded in 3% UltraPureLMP Agarose (Invitrogen, 16520050) dissolved in PBS, onto specimen small dishes; sectioning was performed using Leica vibrating blade microtome VT1200 S (Leica, Wetzlar, Germany), using stainless steel razor blades (Personal Medical, Staunton, VA, USA) under buffered conditions with ice-cold HBSS containing 25 mM glucose (Merck, Darmstadt, Germany), at the following adjustable settings: knife angle: 15°; sectioning speed: 0.4–1 mm/s; oscillation amplitude: 3 mm; step size: 200 lm; retract: 10 lm; continuous stroke.

After sectioning, liver slices (thickness: 200-300 nm) were cultured on top of Nucleopore Track-Etch Membrane (GE Healthcare Life Science, 10417301) to facilitate oxygenation, in 24-well plate in William's E medium (Sigma) supplemented with 25 mM glucose, 20 mM L-Glutamine (Sigma), 1% Pen Strep (Gibco) and 100 mM gemcitabine, at 37°C for 24 hours. Liver slices were washed in PBS and the media was replaced with William's E medium, supplemented with glucose, Pen Strep and with or without rGas6 at a concentration of 400 ng/ml, incubated at 37°C for 24 hours. During the two-day culture the media was replaced every 24 hours.

## **RT-qPCR**

Total RNA purification was performed with the RNeasy Kit (Qiagen), and cDNA was generated using the QuantiTect Reverse Transcription Kit (Qiagen) according to manufacturer's instructions. 500 ng of total RNA was used to generate cDNA. gPCR was performed using 5 x HOT FIREPol EvaGreen gPCR Mix Plus (ROX; Solis Biodyne) on an MX3005P instrument (Stratagene). Three-step amplification was performed (95°C for 15 seconds, 60°C for 20 seconds, and 72°C for 30 seconds) for 45 cycles. Relative expression levels were normalized to Gapdh expression according to the formula: 2^-(Ct gene of interest – Ct Gapdh). Fold increase in expression levels was calculated by the comparative Ct method: 2<sup>^</sup> (-ddCt). The following QuantiTect Primers Assays were used to assess mRNA levels: Mm-Gapdh (Mm Gapdh 3 SG; QT01658692), Mm-Cxcl-1 (Mm Cxcl1 1; QT00115647), Mm-Cxcl2 (Mm Cxcl2 1; QT00113253), Mm-Cxcl-5 (Mm Cxcl5 2; QT01658146), Mm-Gas6 (Mm\_Gas6\_1\_SG; QT00101332), Hs-Gapdh (Hs\_Gapdh\_1\_SG; QT00079247), Hs-Gas6 (Hs\_Gas6\_1\_SG; QT00049126) were purchased from Qiagen. Hs-Cxcl1 (Hs\_Cxcl1\_Fw 5'-AGCTCTTCCGCTCCTCA), Hs-Cxcl2 (Hs\_Cxcl2\_Fw 5'-GAAAGCTTGTCTCAACCCG), 5'-Hs-Cxcl5 (Hs Cxcl5 Fw 5' CCAGTAGTTAGCTTTCTTCCTGGAT), Hs-Cxcl8 (Hs Cxcl8 Fw CCTTCCTGATTTCTGCAGCTC) were purchased from Sigma-Aldrich.

# Immunohistochemistry analysis

Deparaffinization and antigen retrieval were performed using an automated DAKO PT-link. Paraffin-embedded human and mouse liver metastatic sections were immunostained using the DAKO envision system-HRP. Tissue sections were incubated overnight at 4°C with primary antibodies: Cytokeratin 19 (Abcam, ab53119, 1:100); MUC-1 (Abcam, ab89492, 1:100); Ki67 (Abcam, ab15580, 1:1000); CC3 (Cell Signalling, #9661, 1:300); P-AXL (R&D System, AF2228); MPO (Abcam, ab9535, 1:100); YM-1 (StemCell Technology, 60130, 1:200); NKp46 (R&D System, AF2225). Secondary HRP-conjugated antibodies were incubated for 30 minutes at room temperature. Staining was developed using diaminobenzidine and counterstained with haematoxylin. Tumour cells in the liver were identified by their characteristic formation of ductal structures during metastatic growth.

# Immunofluorescence analysis

Murine liver tissues were fixed using a sucrose gradient method to preserve the zsGreen fluorescence. Briefly, livers were fixed in 4% Formaldehyde + 10% sucrose in PBS for 4 hours and then transferred to 20% sucrose in PBS for 8-10 hours. Tissues were transferred into 30% sucrose for an additional 8-10 hours, embedded in OCT medium and stored at - 80°C.

For immunofluorescence staining, 5 mm liver sections were permeabilized by 0.1% TritonX-100 (Sigma Aldrich) for 10 minutes. Unspecific bindings were prevented by using PBS + 8% normal goat serum for 1 hour at room temperature. Tissue sections were incubated overnight at 4°C with the following antibodies: CD8 (BioLegend, 100701, 1:400); CC3 (Cell Signalling Technologies, 9661); F4/80 (BioLegend, 123102, 1:100); iNOS (Abcam, ab15323, 1:50); MUC-1 (Abcam, ab89492, 1:100); Ki67 (Abcam, ab15580, 1:1000); MPO (Abcam, ab9535, 1:50). The next day, tissue sections were washed in PBS and stained with the secondary antibody goat anti-rabbit conjugated to anti-mouse IgG Alexa Fluor(AF)-488 (Abcam, ab150105, 1:200); anti-rabbit IgGAF-488 (Abcam, ab150077, 1:300); anti-rat IgG AF-594 (Abcam, ab150160, 1/300); anti-rabbit IgG AF-594 (Abcam, ab150080, 1/200) and DAPI (Life Technologies, 1:500) for 1 hour at room temperature. Luc/zsGreen-transfected cells were detected by their intrinsic signal. For CD8 staining VECTASTAIN® Elite® ABC-HRP Kit (Peroxidase, Rat IgG) (Vector laboratories) was used. TUNEL staining was performed according to manufacturer's instruction (Thermofisher, 15313038). Sections were finally mounted using Dako Fluorescent Mounting Medium. All tissue sections were imaged using an Axio Observer Light Microscope with the Apotome.2 (Zeiss) and quantified using the Zen Software (Zeiss).

## Statistical analysis

Statistical significance (analysed with GraphPad Prism v5 or v8 software) was determined using two-tailed unpaired Student's t tests when comparing differences between two experimental groups. Unless otherwise stated, for multiple comparisons, one-way ANOVA with Bonferroni was used for all experiments with more than two groups. For Figure 2D, E one-way ANOVA with Dunnett's *post hoc* testing was performed. For Figures 4G, 4J, 8D and Supplementary Figure S5F, N two-way ANOVA with Bonferroni was used. The Log-rank (Mantel-Cox) test was used to assess differences in survival for the metastatic survival study under cytotoxic treatment. A *P*-value < 0.05 was considered statistically significant and *P*-values are indicated in the figures using asterisks: \* *P*<0.05; \*\* *P*<0.01; \*\*\*\* *P*<0.001; \*\*\*\*

Supplementary Table S1. Top upregulated genes GO extracellular.

Supplementary Table S2. Complete gene list GO extracellular.

Supplementary Table S3. Haematology PDAC patients.

# SUPPLEMENTARY METHODS REFERENCES

1 Hingorani SR, Wang L, Multani AS, Combs C, Deramaudt TB, Hruban RH, *et al.* Trp53R172H and KrasG12D cooperate to promote chromosomal instability and widely metastatic pancreatic ductal adenocarcinoma in mice. Cancer Cell 2005;**7**:469-83.

2 Nielsen SR, Quaranta V, Linford A, Emeagi P, Rainer C, Santos A, *et al.* Macrophagesecreted granulin supports pancreatic cancer metastasis by inducing liver fibrosis. Nat Cell Biol 2016;**18**:549-60.

3 D'Costa Z, Jones K, Azad A, van Stiphout R, Lim SY, Gomes AL, *et al.* Gemcitabine-Induced TIMP1 Attenuates Therapy Response and Promotes Tumor Growth and Liver Metastasis in Pancreatic Cancer. Cancer Res 2017;**77**:5952-62.

4 Quaranta V, Rainer C, Nielsen SR, Raymant ML, Ahmed MS, Engle DD, *et al.* Macrophage-Derived Granulin Drives Resistance to Immune Checkpoint Inhibition in Metastatic Pancreatic Cancer. Cancer Res 2018;**78**:4253-69.

5 Finck R, Simonds EF, Jager A, Krishnaswamy S, Sachs K, Fantl W, *et al.* Normalization of mass cytometry data with bead standards. Cytometry A 2013;**83**:483-94.

6 Dobin A, Davis CA, Schlesinger F, Drenkow J, Zaleski C, Jha S, *et al.* STAR: ultrafast universal RNA-seq aligner. Bioinformatics 2013;**29**:15-21.

7 https://www.bioinformatics.babraham.ac.uk/projects/fastqc/.

8 Love MI, Huber W, Anders S. Moderated estimation of fold change and dispersion for RNAseq data with DESeq2. Genome Biol 2014;**15**:550. 9 de Graaf IA, Olinga P, de Jager MH, Merema MT, de Kanter R, van de Kerkhof EG, *et al.* Preparation and incubation of precision-cut liver and intestinal slices for application in drug metabolism and toxicity studies. Nat Protoc 2010;**5**:1540-51.

10 Zimmermann M, Lampe J, Lange S, Smirnow I, Konigsrainer A, Hann-von-Weyhern C, *et al.* Improved reproducibility in preparing precision-cut liver tissue slices. Cytotechnology 2009;**61**:145-52.

# SUPPLEMENTARY FIGURES



**Supplementary Figure S1** 

A Beads exclusion Unable CD45 Unable CD45 Unable CD45 Unable CD45 Unable CD45 Unable CD45 Unable CD45	B220 CD11b Lv6G CD3 KK1.1 NK1.1	CD19 MHC II Ly6C TCRb NK cells	B cells B cells: CD20°CCD19°MHC IIPS F480 CD11c	DeceMHC IIeco MHC IIecoF4/80eco 1beceLy6C <sup>tout</sup> F4/80ecoMHC IIeco cocLy6C <sup>tout</sup> F4/80ecoMHC IIeco F4/80eco
	tSNE1	NK cells: NK1.1 <sup>pc</sup>	Median Low High	



Supplementary Figure S2



**Supplementary Figure S3** 







**Supplementary Figure S6** 

![](_page_44_Figure_3.jpeg)

**Supplementary Figure S7** 

![](_page_45_Figure_3.jpeg)

**Supplementary Figure S8** 

![](_page_46_Figure_3.jpeg)

**Supplementary Figure S9**