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# IL-6 and PD-L1 antibody blockade combination therapy reduces tumor progression in murine models of pancreatic cancer

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# Abstract

**Objective**—Limited efficacy of immune checkpoint inhibitors in pancreatic ductal adenocarcinoma (PDAC) has prompted investigation into combination therapy. We hypothesized that IL-6 blockade would modulate immunologic features of PDAC and enhance the efficacy of anti-PD-L1 checkpoint inhibitor therapy.

**Design**—Transcription profiles and IL-6 secretion from primary patient stellate cells (PSC) were determined via nanostring analysis and immunohistochemistry, respectively. *In vivo* efficacy and mechanistic studies were conducted with antibodies (Ab) targeting IL-6, PD-L1, CD4, or CD8 in subcutaneous or orthotopic models using Panc02, MT5, or KPC-luc cell lines; and the aggressive, genetically engineered PDAC model [Kras<sup>LSL-G12D</sup>, Trp53<sup>LSL-R270H</sup>, Pdx1.cre, Brca2<sup>F/F</sup> (KPC-

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Brca2 mice)]. Systemic and local changes in immunophenotype were measured by flow cytometry or IHC as appropriate.

**Results**—Patient-derived PSC (n=12) demonstrated prominent IL-6 expression, which was localized to stroma of tumors as confirmed via immunohistochemistry. Combined IL-6 and PD-L1 blockade elicited efficacy in in mice bearing subcutaneous MT5 (p<0.02) and Panc02 tumors (p=0.046) which was accompanied by increased intratumoral effector T lymphocytes (CD62L<sup>-</sup> CD44<sup>-</sup>). Administration of CD8-depleting, but not CD4-depleting Ab abrogated the efficacy of combined IL-6 and PD-L1 blockade in mice bearing Panc02 tumors (p=0.0016). This treatment combination also elicited significant anti-tumor activity in mice bearing orthotopic KPC-luc tumors and limited tumor progression in KPC-Brca2 mice (p<0.001). Histologic analysis revealed increased T cell infiltration and reduced  $\alpha$ -SMA<sup>+</sup> cells in tumors from multiple models. Finally, IL-6 and PD-L1 blockade increased overall survival in KPC-Brca2 mice compared to isotype controls (p=0.0012).

**Conclusions**—These pre-clinical results indicate that targeted inhibition of IL-6 may enhance the efficacy of anti-PD-L1 in PDAC.

### Keywords

pancreatic cancer; IL-6; PD-L1

# Introduction

Pancreatic ductal adenocarcinoma (PDAC) is expected to surpass breast and colon cancer, becoming the second leading cause of cancer-related deaths by the year 2030.<sup>1</sup> One of the major difficulties with PDAC is its clinical silence. Typically the disease only becomes apparent after the tumor invades surrounding tissues or metastasizes to distant organs.<sup>2</sup> For many years, the current standard of care for most advanced PDAC patients has been gemcitabine-based regimens including combined therapy with nab-paclitaxel (Abraxane)<sup>3</sup>, or aggressive chemotherapy (e.g. FOLFIRNOX) as a strategy to de-bulk the tumor, and improve candidacy for surgery. Regardless, these advances may rightfully be classified as only incremental, and justify further research to identify novel strategies with potential for long term clinical responses and cures for this devastating malignancy.

Signaling downstream of IL-6 is important in PDAC genesis and progression.<sup>4, 5</sup> This pleiotropic cytokine binds membrane receptor complexes containing the common signal transducing receptor chain GP130 (glycoprotein 130),<sup>6</sup> thereby initiating a complex series of signaling events that include the JAK/STAT, MAPK and PI3K pathways.<sup>7, 8</sup> In particular, STAT3 is activated via phosphorylation at Tyr<sup>705</sup> in most human PDAC specimens, and cooperates with activated *Kras* to drive initiation and progression of PDAC in murine models.<sup>9, 10</sup> The IL-6/STAT3 axis can simultaneously promote the expansion of immunosuppressive cells or alter the balance of T cell subsets. Among the most notable of these subsets are myeloid-derived suppressor cells (MDSCs) and T regulatory cells (T regs), given their prominent expansion and role as poor prognostic indicators in patients with advanced GI cancer.<sup>11–13</sup> Interestingly, data from our group and others point to the pancreatic stroma as one likely source of IL–6. This cytokine is produced in abundance by

components of the stroma including pancreatic stellate cells (PSC) and tumor associated myeloid cells.<sup>5, 14</sup>. In this manner, IL-6 can cooperate with other cytokines, either systemically or in the tumor microenvironment, to further amplify immune changes in patients. Recent studies using an inducible *Kras*-mediated PDAC murine model also showed that IL-6 was instrumental for PDAC progression.<sup>4, 15</sup> In fact, lack of IL-6 completely ablated cancer progression, even in the presence of oncogenic *Kras*.<sup>4</sup> In agreement with these data, recent results from our group emphasize the importance of systemic IL-6 in PDAC patients.<sup>16</sup> We analyzed plasma from untreated patients with metastatic or non-resectable PDAC and found that IL-6, IL-10 and MCP-1 were significantly associated with overall survival, with IL-6 having a strong inverse relationship.<sup>16</sup>

There remains optimism that the dramatic efficacy of immunotherapy in other cancers may be realized in PDAC patients. Indeed, blockade of T cell checkpoint receptors with neutralizing antibodies (Ab) has emerged as a promising immunotherapeutic approach.<sup>17, 18</sup> Programmed death-1-ligand 1 (PD-L1), also known as B7-H1, is a cell surface protein and one of two ligands for program death receptor 1 (PD-1), a costimulatory molecule that negatively regulates T cell responses.<sup>19, 20</sup> Ligation of PD-L1 on cancer cells to PD-1 expressed on T cells suppresses T cell activation and proliferation, and can induce apoptosis. Intratumoral expression of PD-L1 correlates with poor prognosis in patients with PDAC.<sup>17, 21</sup> Indeed, increased PD-L1 expression by cancer cells or the stroma is a fundamental escape mechanism from host immunity.

In theory, IL-6 blockade may modulate immunologic features of PDAC in both the systemic circulation and the tumor microenvironment, thereby enhancing antitumor activity of anti-PD-L1. IL-6 blocking Ab is an attractive component of combination therapy regimens for PDAC for a number of reasons. First, IL-6 targeted Ab are FDA-approved for other indications and could easily be re-purposed for oncology. Second, data from our group and others suggest that IL-6 is a major factor produced by multiple cellular compartments to promote immune suppression and PDAC progression.<sup>17, 22–24</sup>. Finally, blocking IL–6 may tune the phenotypic properties of both myeloid and T cell compartments of a tumor bearing host to be more amenable to efficacy from immunotherapy regimens.

This pre-clinical study tested the hypothesis that IL-6 blockade would enhance the efficacy of anti-PD-L1 checkpoint inhibitor therapy in preclinical models of PDAC. A significant benefit was observed from this therapy in in the subcutaneous MT-5 and Panc02 models, an orthotopic KPC luciferase model, and an aggressive genetically engineered mouse model (GEMM) of PDAC. Mechanistic studies revealed increased pancreatic effector T cell infiltration, and efficacy dependent on CD8<sup>+</sup> T cells. These results indicate that targeting IL-6 may enhance the efficacy of Ab targeting PD-L1 in PDAC by improving the phenotypic properties and infiltration of T cells.

# **Experimental Methods**

### **Cell lines and reagents**

The murine pancreatic cancer cell line Panc02 was purchased from the ATCC and cultured in RPMI (Gibco) with 10% FBS, 10 mM L-glutamine, and antibiotics. Murine MT5

(Kras<sup>LSL-G12D</sup>, Trp53<sup>LSL-R270H</sup>, Pdx1-cre) pancreatic cells were a kind gift from Dr. Tuveson (Cold Spring Harbor Laboratory, Cold Spring Harbor, NY) and cultured in RPMI (Gibco) with 10% FBS, 10 mM L-glutamine, and antibiotics. KPC-luc cells for orthotopic experiments were derived from KPC mice (Kras<sup>LSL-G12D</sup>, Trp53<sup>-/-</sup>, PDX-1-Cre) and transfected with enhanced firefly luciferase as previously described.<sup>25</sup> Murine isotype controls (clone 3E5.2H12), anti-IL-6R (clone BP-5875), and anti-PD-L1 blocking Ab were obtained from either Genentech, Inc. (San Francisco, CA) for *in vivo* studies in the KPC-Brca2 murine model. Murine antibodies to IL-6 (Clone MP5-20F3), PD-L1 (Clone 10F. 9G2), or isotype controls (Clones LTF-2 and HRPN) were purchased from BioXcell (West Lebanon, NH) for *in vivo* studies using the MT-5, Panc02, and KPC-luc cell lines.

### Murine models of pancreatic cancer

KPC-Brca2 mice were generated by interbreeding *Brca2<sup>flox2/flox2</sup>; Kras<sup>LSL-G12D/+</sup>* with *Brca2<sup>flox2/flox2</sup>; Trp53<sup>LSL-R270H/+</sup>; Pdx1-cre* animals.<sup>26</sup> The mouse strains *p53<sup>LSL-R270H</sup>* (strain number 01XM3), *Kras<sup>LSL-G12D</sup>* (strain number 01XJ6), and *Pdx1-cre* (strain number 01XL5) were acquired from the National Cancer Institute (NCI) Frederick Mouse Repository. All transgenic mice generated in this study were maintained on a mixed 129/B6 genetic background. All studies involving MT5, Panc02, KPC-luc tumors utilized syngeneic, female C57BL/6 mice, 5–6 weeks of age.

### In vivo efficacy studies

KPC-Brca2 mice (5 weeks of age) were treated with isotype controls, anti-IL-6R and/or anti-PD-L1 Ab (Genentech) at a dose of 20µg/mouse, 3 times each week (Monday, Wednesday, and Friday). Following 2 weeks of treatment, animals were euthanized via CO<sub>2</sub> asphyxiation, followed by cardiac puncture. Plasma, splenocytes and tumor tissue were collected for further analysis. Pathology was assessed in H&E stained slides to determine the differentiation state of tissue as pancreatic intraepithelial neoplasia (PanIN) 1A, PanIN 1B, PanIN 2, or PDAC. For studies using MT5 and Panc02 tumors,  $1 \times 10^6$  or  $3 \times 10^5$  cells, respectively were injected subcutaneously in the flank of C57BL/6 mice 3 times each week with 20µg/mouse of isotype, anti-IL-6 or anti-PD-L1 Abs (BioXCell) Ab treatment starting once tumors reached 50-100mm<sup>3</sup> volume. For orthotopic studies, C57BL/6 mice were injected with  $1 \times 10^6$  KPC-luc (luciferase expressing) cells in Matrigel (BD Biosciences) in the tail of the pancreas. Tumor growth was analyzed once a week by bioluminescent imaging and end of study tumor weight was determined immediately post-mortem. Mice were treated 3 times each week with 200 µg/mouse of isotype, anti-IL-6 or anti-PD-L1 Abs (BioXCell). For T cell depletion studies, Ab to deplete CD4 (Clone GK1.5; BioXcell) or CD8 (Clone 2.43; BioXCell) were injected i.p. at 100  $\mu$ g per mouse on days -2, -1, +1, +4, and every other 3 days afterwards until completion of the study as previously described.<sup>27</sup> For survival studies, KPC-Brca2 mice were treated starting at 5 weeks of age with isotype control, IL-6 or PD-L1 Abs as single agents or in combination (200 µg/mouse each Ab, BioXCell) until mice were moribund as determined by IACUC protocol.

#### Pancreatic stellate cell isolation and Nanostring analysis

Tissue from human patient pancreatic tumors undergoing surgical resection at the James Cancer Hospital and Solove Research Institute (Columbus, OH) was obtained under an

Institutional review board-approved protocol following informed consent. Tissue was dissected with a scalpel into 0.5-1mm<sup>3</sup> pieces, then plated in 6-well 10 cm<sup>2</sup> uncoated culture wells in DMEM with 10% FBS and antibiotics and incubated at 37°C. PSC typically grew out of the tissue in 2–3 weeks, and were characterized by morphology and histological analysis of alpha-smooth muscle actin ( $\alpha$ -SMA<sup>+</sup>) staining. PSC were maintained in culture with fresh media added twice weekly for 3 passages and then RNA was collected by Trizol extraction. RNA was analyzed using the nCounter PanCancer Immune Profiling Panel (Nanostring Technologies, Seattle WA).

#### Immunohistochemical (IHC) analysis

Tumor tissue was prospectively obtained at surgery under an Institutional review boardapproved protocol following informed consent and de-identified. Formalin fixed pancreatic tissue from human tissue specimens and *in vivo* experiments in mice were subjected to IHC analysis following staining with Ab against IL-6 (Abcam and eBioscience), pSTAT3 (Catalog 4904; Cell Signaling),  $\alpha$ -SMA (Catalog M0851; Dako), PD-L1 (Catalog NBP1-76769) and CD3 (Catalog A0452; Dako). For  $\alpha$ -SMA quantification, 20× magnification images of pancreata (8–10 images per mouse) were captured using PerkinElmer's Vectra multispectral slide analysis system. inForm software tools were used to quantify positive SMA cells (Fast Red chromogen) within each image (Supplementary Figure 1). For analysis of CD3<sup>+</sup> cells, blinded histological analysis of staining in the pancreas was counted at 40× magnification, with at least 10 fields counted per mouse.

### **Flow Cytometry**

Immunophenotypic analyses of splenoctyes and single cell suspensions from tumors were assessed by flow cytometry. Antibodies to stain for MDSC were CD11b-APC (Clone M1/70; BD Biosciences), Ly6G-FITC (Clone 1A8; BD Biosciences), Ly6C-PE (Clone AL-21; BD Biosciences). For T cell activation markers, cells were stained with Ab specific for CD4-PE-Cy7 (Clone RM4-5; BD Biosciences), CD8-PE-Cy7 (Clone 53-6.7; BD Biosciences), CD62L-PE (Clone MEL-14; BD Biosciences), and CD44-FITC (Clone IM7; Biolegend). To determine Th1 and Th2 phenotypes, cells were stained using fluorochrome-conjugated Ab targeted CXCR3-PE-Cy7 (Clone CXCR13-173; Biolegend), CCR4-PE (Clone 2G12; Biolegend), and CCR6-APC (Clone CK4-L3; BD Biosciences). Cells were incubated on ice for 30 minutes, washed, and fixed in PBS containing 1% formalin for flow cytometric analysis on a LSRII flow cytometer (BD Biosciences).

#### Statistics

For data obtained by Nanostring (Nanostring Technologies, Seattle, WA), positive spike-in controls were first used to normalize assay efficiency. Negative controls were used to access background hybridization, and for filtering out low expression probes. Reference genes were used to normalize across biological samples. Linear mixed models were then used to detect differentially expressed RNAs. A variance smoothing method was used for improving variance estimates in testing.<sup>28</sup> Significance was adjusted by controlling the mean number of false positives.<sup>29</sup> Data obtained by flow cytometry and IHC as well as tumor volumes were log-transformed prior to analysis to meet model assumptions of normality and homoscedasticity. Poisson regression was used to model the number of PanIN3/tumor

lesions in the KPC-Brca2 mice. Tumor volume was modeled over time using mixed-effects regression with fixed effects for group, time and the interaction of the two. Random intercepts and slopes by mouse were included with an unstructured covariance matrix for the random effects. For the T cell depletion experiments examining tumor growth, interaction contrasts were constructed to enable the direct testing of the hypotheses under consideration.<sup>30</sup> For Kaplan-Meier survival curves a log-rank test to determine difference used. The other outcomes were compared using ANOVA. P-values less than 0.05 were considered significant and all analyses were conducted in SAS v9.4 (SAS Institute, Cary, NC) or R v3.2.3 (R Foundation for Statistical Computing, Vienna, Austria).

# Results

#### The pancreatic tumor stroma is a major source of IL-6 in the tumor microenvironment

For these studies, an unsupervised assessment of gene expression was conducted to identify actionable immunologic targets in pancreatic stellate cells, a prominent component of PDAC stroma. Stellate cells derived from surgically resected pancreatic tissue of 10 different patients were isolated and cultured *in vitro* as previously described.<sup>14</sup> After 3 passages, RNA was extracted and subjected to profiling of transcripts relevant to immunologic biomarkers using Nanostring technology (Nanostring PanCancer Immune Profiling Panel). RNA isolated from patient-derived PSC exhibited significantly increased expression of cytokines and chemokines including IL-6, and CCL11 as compared to RNA isolated from normal human pancreatic fibroblasts (Figure 1 and Supplementary Figure 2). Further investigation via IHC analysis confirmed a pattern of staining for IL-6 that localized predominantly in the desmoplastic, stromal regions of pancreatic tumors, with occasional staining in regions containing malignant pancreatic epithelial cells (Figure 2A). This trend was evident in tissue from several (n=13) representative patients (Figure 2B). These data were consistent with our analysis of transcripts in PSC, and suggested that the stromal compartment, is a major source of IL-6 in the pancreatic tumor microenvironment.

# Prominent expression of IL-6/JAK/STAT pathway components and PD-L1 in the pancreatic tumor microenvironment of mouse models of pancreatic cancer

Similar to human PDAC, histologic characterization of pancreata from PDAC orthotopic and genetically engineered mouse models (GEMM) revealed expression of IL-6 in both stroma, and more limited positive staining in foci of some tumors (Figure 3). Consistent with prior published studies, pancreata from GEMM (KPC and KPC-Brca2) recapitulated the histologic features of advanced PDAC observed in human disease. For example, KPC tumors displayed acinar to ductal metaplasia, while the KPC-Brca2 displayed desmoplastic stroma, PanIN, and ductal adenocarcinoma expression similar to observations in human tissue. Pancreata from both the orthotopic and genetic tumor models displayed a fibrotic stroma that both surrounded and intercalated within tumor foci (Figure 3; S, stroma; T, tumor tissue). Of note, robust phosphorylation of STAT3 was detectable in stroma and malignant epithelial cells in the pancreata from both KPC and KPC-Brca2 mice (Figure 3). Consistent with this immunosuppressive microenvironment, tumors expressed high levels of PD-L1, regardless of the animal model.

# Combined blockade of IL-6 and PD-L1 inhibits murine pancreatic tumor growth and increases intratumoral effector T cells *in vivo*

We postulated that concurrent blockade of IL-6 and a potent immune-modulatory stimulus via PD-L1-targeting Ab would circumvent key mechanisms of resistance to T-cell mediated immunity to pancreatic cancer. We utilized murine MT5 tumor cells subcutaneously injected into C57BL/6 mice. This cell line was derived from KPC tumors and has both G12D mutated Kras and R172H Trp53.31 Once tumors were palpable, mice were treated with anti-IL-6, anti-PD-L1 or isotype controls until study endpoint. A significant inhibition of tumor growth was observed in response to the combination therapy as compared to isotype, anti-IL-6 or PD-L1 antibody therapy alone (Figure 4A; p<0.03). To carefully assess immune changes, and confirm our data in an additional PDAC model, the well-characterized, murine Panc02 model was also used, given its frequent use in published pre-clinical immunotherapy studies in PDAC. Similar to the MT5 cell line, Panc02 cells are also syngeneic to immune competent C57BL/6 mice.<sup>32, 33</sup> Once tumors were palpable, mice received treatment until the study endpoint. A significant inhibition of tumor growth was observed in response to combined therapy with anti-IL-6 and anti-PD-L1 Ab as compared to isotype controls (Figure 4B; p=0.0457). Tumors were dissociated at the study endpoint and underwent phenotypic analysis by flow cytometry. These data confirmed a significantly greater percentage of infiltrating CD8<sup>+</sup> T cells in the tumor tissue from mice receiving the combination (Figure 4C; p=0.0006). Further analysis indicated an increased proportion of cells with an effector CD62L<sup>-</sup>CD44<sup>+</sup> phenotype in mice receiving combined blockade of IL-6 and PD-L1 (Figure 4D; p=0.0086 and E; p=0.0024). Further, we observed no difference in the level of other circulating or intratumoral immunosuppressive subsets (Granulocytic and Monocytic MDSC) between any of the treatment groups as compared to isotype controls (Supplementary Figure 3A-C). Importantly, treatment Ab blocking IL-6, PD-L1 or both Ab combined did not result in any overt signs of toxicity as evidenced by weight loss in all animal models tested (data not shown).

# Decreased tumor growth by combined blockade of IL-6 and PD-L1 is CD8<sup>+</sup> T cell dependent

To determine whether the efficacy of combined IL-6 and PD-L1 blockade was T cell dependent, Panc02-bearing mice were administered CD4<sup>+</sup> or CD8<sup>+</sup> depleting Ab during the course of treatment. T cell depletion was confirmed by flow cytometry using splenocytes from mice at the study endpoint (Figure 5A and B). Administration of CD8 (p=0.0016), but not CD4 depleting Ab abrogated the efficacy of combined IL-6 and PD-L1 blockade in mice bearing Panc02 tumors (Figure 5C).

# IL-6 and PD-L1 antibody blockade combination therapy reduces tumor progression in an orthotopic model of PDAC

To further investigate the efficacy of IL-6 blockade combined with PD-L1 inhibition, we utilized KPC-luc cells orthotopically injected into the tail of the pancreas of C57BL/6 mice. This luciferase-expressing cell line was derived from KPC tumors with G12D mutated *Kras* and R172H *Trp53*.<sup>25</sup> Tumor growth was assessed by weekly bioluminescent imaging (Figure 6A and B). Once tumors were detectable by imaging the pancreas, mice were randomized

and treated with anti-IL-6, anti-PD-L1 or isotype control Ab until the study endpoint. We observed a decreased final tumor weight in mice receiving combination therapy as compared to isotype control, anti-IL-6, or PD-L1 antibody therapy alone (Figure 6C; p=0.05530). A decrease in  $\alpha$ -SMA<sup>+</sup> stromal cells was also observed in the pancreata of mice treated with combination therapy as compared to isotype control Ab alone (p=0.0563). Furthermore, the number of CD3<sup>+</sup> T cells was greater in the pancreata from mice treated with antibodies targeting both IL-6 and PD-L1 as compared to those treated with isotype control antibodies alone (p=0.0553).

# Antibodies targeting IL-6R and PD-L1 limit tumor progression in the aggressive KPC-Brca2 GEMM

To further confirm our data in a genetic model that better recapitulates many aspects of patient tumors, we used a highly aggressive, autochthonous model of spontaneously arising PDAC driven by mutant Kras, Trp53 and Brca2 (Figure 7A). Any efficacy in this oncogene driven tumor model would signify a promising approach to carry forward for further evaluation. At 5-6 weeks of age, KPC-Brca2 mice have PDAC with 100% penetrance (Figure 7A). Following two weeks of treatment with anti-IL-6R and/or anti-PD-L1, animals were euthanized and pancreata was isolated for histologic analysis. There was a statistically significant shift in the proportion of low grade PanIN lesions, with fewer PanIN3 or foci of adenocarcinoma in mice treated with antibodies targeting PD-L1 and IL-6R, as compared to animals treated with isotype control Ab (Figure 7A and B; p<0.001). Similar to the subcutaneous Panc02 model, depletion of CD8<sup>+</sup> T cells limited the effect of combination therapy in the KPC-Brca2 GEMM. In fact, we observed a greater proportion of PanIN3/ PDAC lesions in pancreata from CD8<sup>+</sup> depleted mice treated with the combination when compared to mice treated with the combination alone (Figure 7B). Importantly, analysis revealed a significant decrease in the number of  $\alpha$ -SMA<sup>+</sup> stromal cells in pancreata of mice treated with Ab targeting both IL-6R and PD-L1 as compared to control mice (Figure 7C and D; p<0.05). Mice depleted of CD8<sup>+</sup> T cells treated with combination therapy did not display the same decrease of  $\alpha$ -SMA<sup>+</sup> cells in the pancreas compared to mice treated with only the combination therapy (Figure 7D; p=0.0033). Further, treatment with Ab targeting IL-6, PD-L1 or both agents combined did not alter expression of IL-6 (Supplementary Figure 5A) or PD-L1 (Supplementary Figure 5B) in the pancreatic tumor microenvironment.

# Combined blockade of IL-6R and PD-L1 increases T cells with Th1 phenoytpic properties, and CD3<sup>+</sup> cell infiltration into PDAC tumors in KPC-Brca2 mice

To better understand the mechanism by which combined IL-6 and PD-L1 blockade limited tumor progression in the aggressive KPC-Brca2 model, a series of immunophenotypic analyses were conducted. These data revealed a higher percentage of circulating CD4<sup>+</sup> T cells with Th1 phenotypic properties (CXCR3<sup>+</sup>CCR4<sup>-</sup>CCR6<sup>-</sup>) in the spleens of mice treated with Ab targeting IL-6R and PD-L1 combined as compared to controls (Figure 8A; p<0.0001). No changes in CD4<sup>+</sup> T cells with Th2 phenotypic properties were evident in any treatment group as compared to controls (Figure 8B). Importantly, assessment of the tumor microenvironment revealed significant increases in the infiltrating CD3<sup>+</sup> T cells in the pancreata of combination treated mice (Figure 8C–D; p=0.0025). Further analysis revealed no difference in the level of other immunosuppressive subsets (MDSC and Tregs) between

any of the treatment groups as compared to isotype controls (Supplementary Figure 4A–D). These data suggest that this treatment regimen may modulate phenotypic properties of T cells and their access to the tumor microenvironment.

#### Combined blockade of IL-6 and PD-L1 extend overall survival in the KPC-Brca2 GEMM

The impact of this treatment combination on overall survival was evaluated in KPC-Brca2 mice. In this study, mice began long-term treatment at 5 weeks of age with antibodies targeting IL-6 and PD-L1 or isotype controls 3 times weekly until the animals were moribund. A significant prolongation in overall survival (35%; p=0.0012) was observed in mice treated with combined IL-6 and PD-L1 Abs, as compared to isotype control treated animals (Figure 9). Long term exposure to this treatment regimen did not lead to alterations in body weight of these animals, indicating its tolerability during a sustained period of administration (Supplementary Figure 6).

# Discussion

In the present report we demonstrate that combined blockade of IL-6 and PD-L1 elicits efficacy and extends survival therapy in highly aggressive PDAC models. These data extend prior observations regarding IL-6 as a factor limiting the efficacy of immunotherapy in PDAC. Combination therapy skewed T cells to a Th1 phenotype, and increased the number of infiltrating effector cells into the tumor. This is significant, as limited T cell infiltration is considered a barrier for efficacy of immunotherapy in PDAC. Further, we showed that efficacy of combination therapy on tumor progression was dependent on CD8<sup>+</sup> T cells in multiple models. Overall, these pre-clinical data support this combination as a rational, therapeutic approach to elicit improved T cell mediated immune responses to pancreatic cancer.

The next frontier of immunotherapy is discovering ways to enhance the efficacy of checkpoint inhibitors using combination approaches, especially in cancers where little effect is observed in the clinic. Before blindly moving forward with combination approaches, it will be important to prioritize therapies based on empirical data, and the relative importance of targeting each factor in the tumor-bearing host. IL-6 as a target is quite attractive for a number of reasons. For example, as recently reported by our group, higher baseline plasma IL-6 in treatment-naïve PDAC patients correlated with worse overall survival.<sup>16</sup> IL-6 can also promote the differentiation of DCs, macrophages, and MDSC contributing to immunosuppression.<sup>19, 22</sup> Further, IL-6 can lead to greater proliferation, invasiveness, and tumor progression in the pancreatic tumor microenvironment.<sup>16</sup> This report represents an important series of pre-clinical studies to provide the first evidence of this treatment combination working in relevant animal models. The next logistical step for effective translation will be to understand how this drug combination works in the context of prior or concurrent chemotherapy, which will be most relevant in clinical trial scenarios.

IL-6 targeting agents have been utilized in pancreatic cancer but with limited success. This is potentially because IL-6 blockade has been attempted as a single agent, or in combination with chemotherapy in heavily pre-treated patients.<sup>34, 35</sup> In all of these studies, there was no potent immunotherapeutic treatment that was given concurrently with blockade of IL-6.

Others have shown that neutralization of IL-6 also suppresses signaling through the phosphoinositide 3-kinase/AKT pathway.<sup>36</sup> A recent Phase-I study combining tocilizumab (anti-IL-6) and IFN-a2b in ovarian cancer showed feasibility/safety and a greater percentage of activated T cells.<sup>37</sup> Thus, blockade of IL-6 could lead to potentially more efficacious results than targeting a single pathway (JAK/STAT, MEK, AKT) as it could simultaneously limit other pathways driven by aberrant IL-6 expression.

Our group and others have reported that tumors, myeloid cells, and stromal cells can secrete IL-6, fueling the tumor microenvironment and immunosuppression in PDAC. It is known that PDAC is profoundly immune suppressive, and clearly more data on clinical relevance of the stroma and its soluble factors, including IL-6 are needed. In a model of MCA205 cells transfected with OVA, Gr-1<sup>+</sup> MDSC were the source of IL-6 in tumor bearing mice. Blockade of IL-6 led to increased differentiation of tumor-Ag specific effector T helper cells (Th1).<sup>24</sup> In this report, IL-6 in stellate cells derived from PDAC stroma was a key factor that emerged in an unsupervised analysis of patient PSC at the RNA level and was confirmed via IHC. This was consistent with our prior in vitro data showing that patient-derived PSC secrete IL-6 that promotes the differentiation of MDSC.<sup>20</sup> Other studies have clearly shown that IL-6 can arise from tumor derived macrophages, and may play a role via STAT3 in mediating immune suppression in the tumor microenvironment.<sup>5</sup> These factors complement other cytokines such as GM-CSF, which is produced by tumor cells (and perhaps others) to fuel an immunosuppressive microenvironment.<sup>38</sup> These data indicate that IL-6 represents an actionable therapeutic target that may modulate the immune architecture in PDAC tumors and can be manipulated to limit immune suppression.

It is likely that the PDAC stroma amplifies immunosuppression via IL-6-induced signaling, and limits the efficacy of immunotherapy. For instance, numerous reports suggest that a reciprocal relationship exists between tumor and stromal cells, leading to enhanced survival and growth of PDAC.<sup>39</sup> Among these are studies demonstrating that fibroblast activating protein (FAP<sup>+</sup>) stromal cells maintain the stroma and suppress anti-tumor immunity.<sup>40–42</sup> Other studies, using targeted disruption of sonic hedgehog signaling in the stroma led to accelerated PDAC progression.<sup>43, 44</sup> These data provided insight into prior clinical studies using the hedgehog pathway inhibitor Saridegib which were halted early due to patients progressing faster than those on the control arm. Similarly, genetic ablation of SMA<sup>+</sup> stromal cells led to accelerated metastasis and disease progression in a mutant *KRas*-driven model of PDAC.<sup>45</sup> Despite these data, it was quite interesting to note that the response to immunotherapy with anti-CTLA4 Ab was enhanced in animals lacking SMA<sup>+</sup> stroma cells. In contrast, these animals did not derive benefit from chemotherapy with geneticabine.<sup>45</sup>

Subsequent studies should investigate further how IL-6 blockade enhances checkpoint immunotherapy. IL-6 could act to promote expansion of MDSC, although no alterations were observed in MDSC levels in tumor and/or spleen in either the genetic or subcutaneous murine models (Supplementary Figures 3 and 4). MDSC and macrophages are other potential sources of IL-6, and blockade could also render these cells less immunosuppressive, leading to enhanced efficacy of checkpoint inhibition. IL-6 could act at the level of antigen presenting cells, by down-regulating IL-12 production, and subsequently the Th1 immune response.<sup>24, 46</sup> Another possibility is that IL-6 blockade could

downregulate SOCS expression in T-cells, and restore the ability of IFN-gamma or other cytokines to act and promote Th1 differentiation.<sup>47</sup> Although this is less likely given the redundancy in cytokines that might compensate and upregulate SOCS expression, it remains a possibility.

One final interesting and important observation was that anti-IL-6 and anti-PD-L1 combination increased T cell infiltration into tumors. T cell infiltration into tumors is a major issue that impacts the ability of a patient to respond to checkpoint inhibitors.<sup>48, 49</sup> This regimen clearly adjusts the phenotypic properties of T cells that infiltrate tumors, and may make a subject more likely to respond to immunotherapy. Blockade of IL-6 in combination with checkpoint therapy could improve T cell trafficking, and change the profile of T cells in the tumor. We believe this combination may be a broadly applicable therapeutic strategy for multiple types of tumors that are driven by IL-6. Given the multiple facets of IL-6 in regulating pancreatic carcinogenesis and immune suppression, this represents a reasonable target with a rapid path to clinical translation since multiple IL-6 blocking Ab are already FDA approved for other indications.

# **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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# Abbreviations

PDAC	pancreatic ductal adenocarcinoma
IL-6	interleukin-6
PD-L1	Programmed death-1-ligand 1
PSC	pancreatic stellate cell
GEMM	genetically engineered mouse model
JAK	Janus kinases
STAT	signal transducer and activator of transcription

PanIN	pancreatic intraepithelial lesions
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Treg T regulatory

MDSC myeloid derived suppressor cell

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### Significance of this study

### What is already known on this subject?

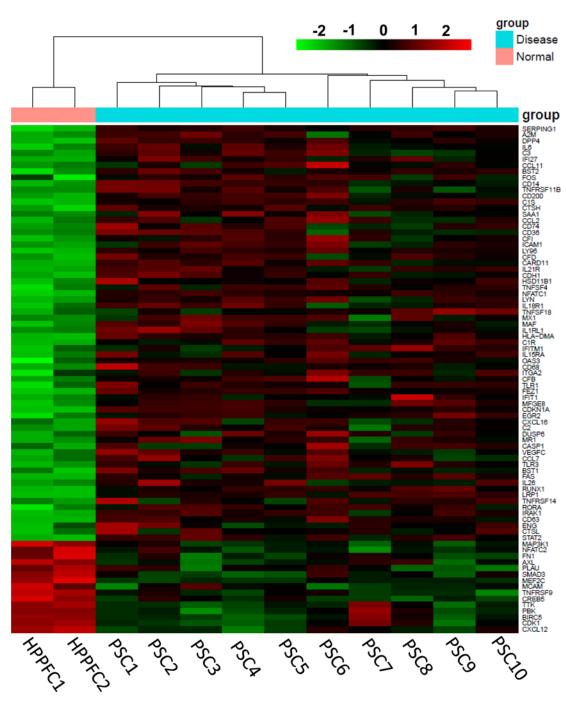
- Checkpoint blockade immunotherapy has limited efficacy in pancreatic cancer patients.
- The pro-inflammatory cytokine, IL-6 is highly elevated in pancreatic cancer patients and is a prognostic factor for outcome.

# What are the new findings?

- Pancreatic stellate cells are a major source of IL-6 in the pancreatic tumor microenvironment.
- Blockade of IL-6 and PD-L1 limits tumor progression and enhances overall survival in aggressive murine models of pancreatic cancer.
- Combination therapy increased circulating cells with Th1 phenotypic characteristics and intratumoral effector T cells.
- Combined blockade of IL-6 and PD-L1 immunotherapy elicits efficacy in a CD8<sup>+</sup> T cell-dependent manner.

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

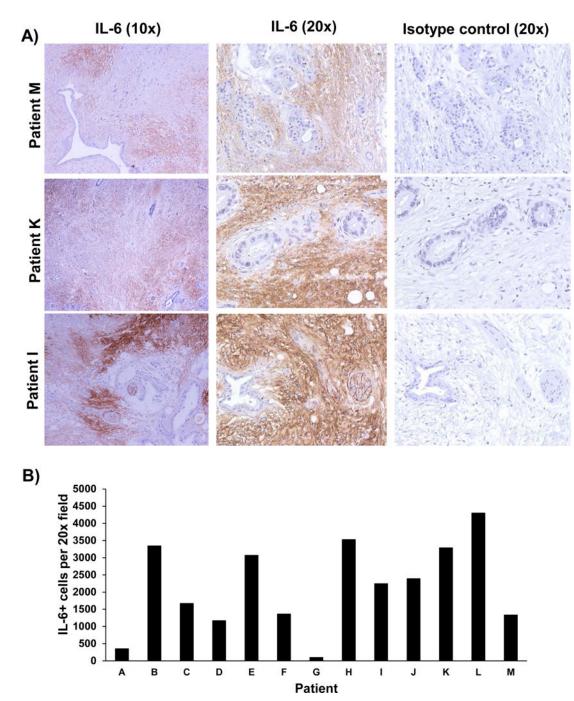
- These preclinical results elucidate a novel combination therapy that could be translated to the clinic.
- Our results suggest that targeting IL-6 could improve pancreatic cancer patient responses to checkpoint blockade immunotherapy.



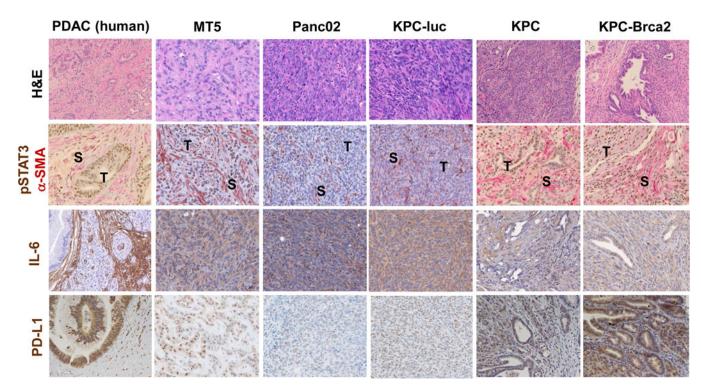
#### Figure 1. Profile of mRNA transcripts from patient-derived PSC

RNA was isolated from 10 patient-derived pancreatic cancer stellate cells and analyzed utilizing the Nanostring nCounter PanCan Immune Profiling Panel. Data are expressed as the fold change in expression as compared to a normal human pancreatic fibroblast cell line and relative to several housekeeping genes. Genes are ranked from highest fold-change to lowest. Heat map with hierarchical clustering for genes with at least 2 fold change up or down with a p<0.01 cutoff. Significantly higher expression is shown in red and lower expression in green.

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**Figure 2. IL-6 expression is enriched in the stromal compartments of human PDAC tissue** Human PDAC tumors (among n=13 patients) stained for IL-6 expression by immunohistochemistry. **A)** IL-6 (Brown Chromogen) IHC staining from three representative patient samples is displayed at 10× and 20× magnification. Tissue sections from each patient were stained with an isotype control Ab to account for background. **B**) Number of IL-6 positive cells per field was quantified.



# Figure 3. PDAC GEMM recapitulate IL-6 pathway activation and PD-L1 expression observed in human disease

IHC analysis of representative H&E, pSTAT3 (brown)/ $\alpha$ –SMA (pink), IL-6 (brown), and PD-L1 (brown) staining on tissue sections from human pancreatic cancer, orthotopic (KPC-luc), subcutaneous (MT5, Panc02), or pancreata from the following GEMM: KPC (Kras<sup>LSL-G12D</sup>, Trp53<sup>LSL-R270H</sup>, Pdx1-cre), and KPC-Brca2 (Kras<sup>LSL-G12D</sup>, Trp53<sup>LSL-R270H</sup>, Pdx1-cre). All images are presented at 40× magnification.

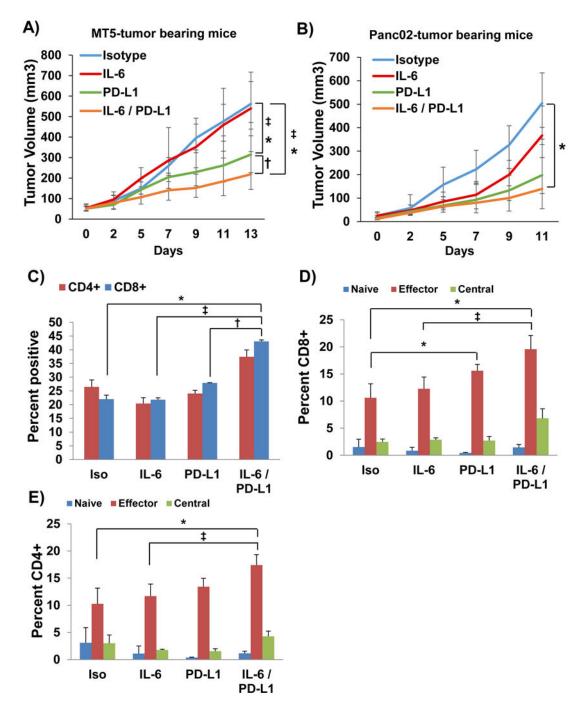
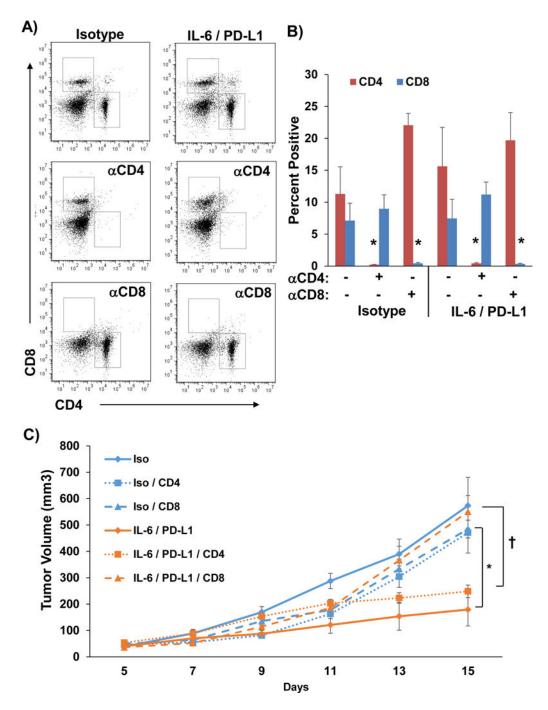
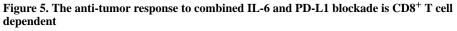


Figure 4. IL-6 and PD-L1 antibody blockade combination therapy decreases PDAC tumor progression and increases the percentage of intratumoral effector T cells A) MT5 or B) Panc02 murine pancreatic tumor cells were subcutaneously injected into C57BL/6 mice with treatment beginning when tumors reached 50–100mm<sup>3</sup>. Mice were treated with 200µg (intraperitoneal injection 3 times/week) with isotype control, anti-IL-6 and/or anti-PD-L1 antibodies (n=5–6 mice/group) until mice met pre-specified IACUC-approved early removal criteria. Geometric means ± SD; \*p<0.01 compared to Isotype; <sup>†</sup>p<0.03 compared to PD-L1; <sup>‡</sup>p<0.05 compared to IL-6. Panc02 tumors were

dissociated using Collagenase II and the Miltenyi Biotec gentleMACS dissociator to obtain a single cell suspension. Cells were stained and analyzed by flow cytometry for C) CD4<sup>+</sup>, CD8<sup>+</sup> T cells and activation markers for different D) CD4<sup>+</sup> and E) CD8<sup>+</sup> T cell subsets (n=3/group). Naïve (CD62L<sup>+</sup>CD44<sup>-</sup>), Effector Memory (CD62L<sup>-</sup>CD44<sup>+</sup>), and Central Memory (CD62L<sup>+</sup>CD44<sup>+</sup>). Means  $\pm$  SD; \*p<0.002 compared to Isotype; <sup>†</sup>p<0.01 compared to PD-L1; <sup>‡</sup>p<0.02 compared to IL-6.





Panc02 murine pancreatic tumor cells were subcutaneously injected into C57BL/6 mice with treatment beginning when tumors reached 50–100mm<sup>3</sup>. Mice were also depleted of either CD4<sup>+</sup> or CD8<sup>+</sup> T cells by injecting depletion antibodies on days 5, 6, 8, 11, and 14. **A**) Representative flow cytometry and **B**) quantification of CD4 and CD8 staining of splenocytes to confirm depletion. Means  $\pm$  SD; \*p<0.001. **C**) On day 7, mice were treated with 200 µg (intraperitoneal injection 3 times/week) of isotype controls or anti-IL-6 and

anti-PD-L1 antibodies combined (n=5 mice/group) until mice met pre-specified IACUCapproved early removal criteria. Means  $\pm$  SD; \*p<0.001 compared to isotype; <sup>†</sup>p<0.002 CD8 depletion vs CD4 depletion.

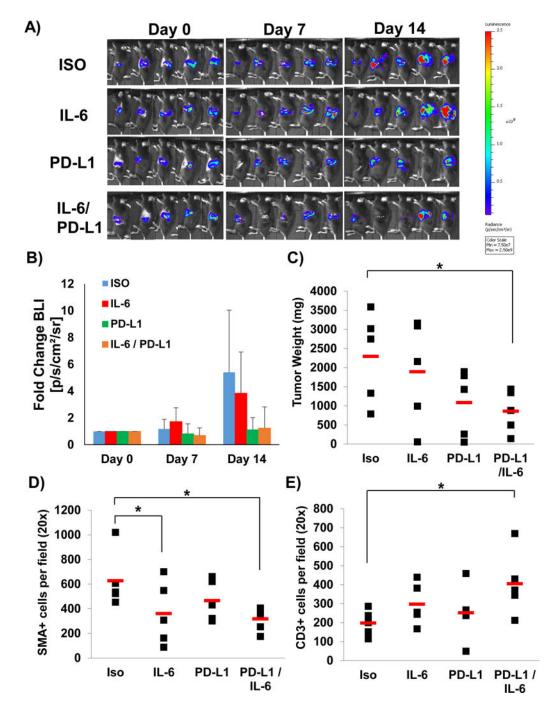
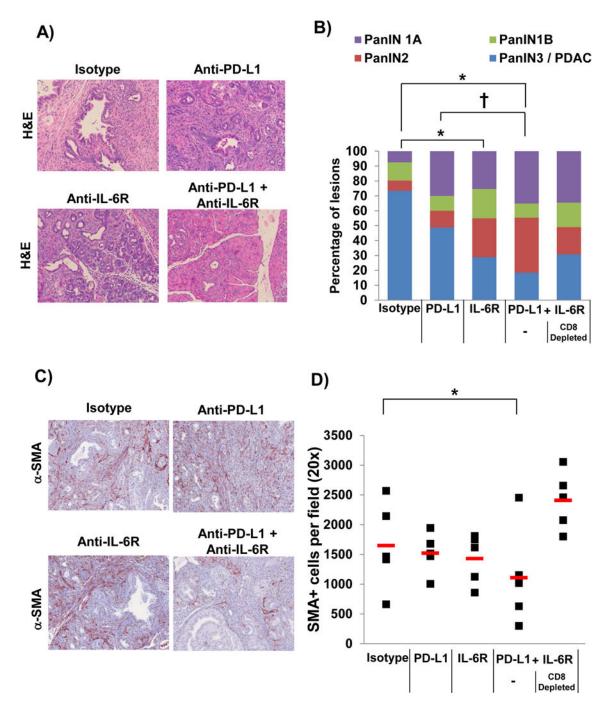
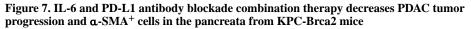


Figure 6. IL-6 and PD-L1 antibody combination blockade reduces tumor growth in an orthotopic model of pancreatic cancer

C57BL/6 mice were orthotopically injected with luciferase expressing KPC cells (KPC-luc) in matrigel and **A**) tumor growth assessed and **B**) quantified by bioluminescent imaging weekly. **C**) Tumor weights at completion of study. Means  $\pm$  SD; \*p=0.05530 compared to isotype. Pancreatic tissue was stained for **D**)  $\alpha$ -SMA<sup>+</sup> stromal cells (Means  $\pm$  SD; \*p=0.0563 compared to Isotype) and E) CD3<sup>+</sup> T cells (Means  $\pm$  SD; \*p=0.0553 compared to Isotype) by IHC quantified at 20× magnification.

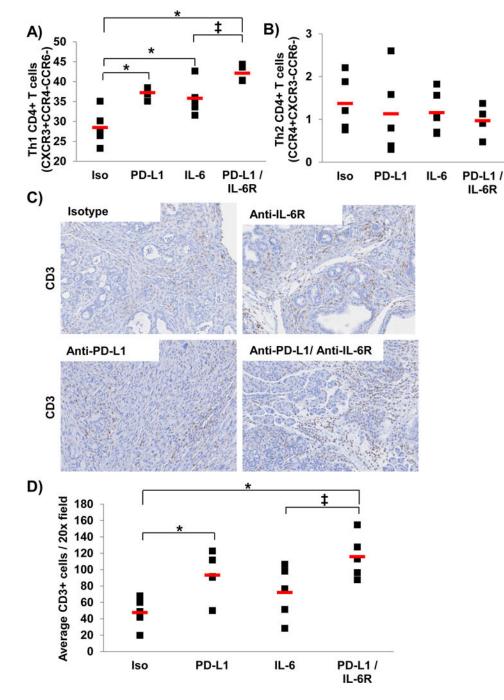
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KPC-Brca2 mice were treated at 5–6 weeks of age with 200  $\mu$ g (intraperitoneal injection 3 times/week) of isotype control, anti-IL-6R and/or anti-PD-L1 antibodies for 2 weeks (n=5 mice/group). A cohort of mice treated with anti-IL-6R and PD-L1 antibodies were also depleted for CD8<sup>+</sup> T cells. **A**) Representative H&E staining of pancreata at 20× magnification and **B**) quantification of the pathology. Means ± SD; \*p<0.005 compared to Isotype; <sup>†</sup>p=0.01) compared to PD-L1; **C**) Pancreatic tissue was stained for  $\alpha$ -SMA<sup>+</sup>

stromal cells (red) by IHC and **D**) quantified at 20× magnification. Means  $\pm$  SD; \*p=0.0545 compared to Isotype.

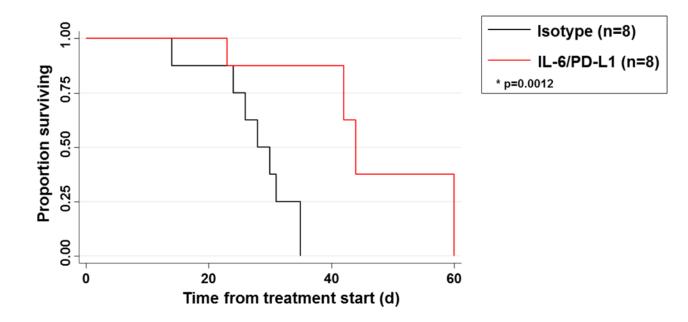


**Figure 8. Higher percentage of circulating T cells with Th1 phenotypic properties and intratumoral CD3<sup>+</sup> cells in mice treated with IL-6 and PD-L1 antibody blockade** KPC-Brca2 mice were treated at 5–6 weeks of age with 200 μg (intraperitoneal injection 3 times/week) with isotype control, anti-IL-6R and/or anti-PD-L1 antibodies for 2 weeks (n=5 mice/group). Splenocytes were stained and analyzed by flow cytometry for **A**) T cells with Th1 phenotypic properties (CD4<sup>+</sup>CXCR4<sup>+</sup>CCR4<sup>-</sup>CCR6<sup>-</sup>) or **B**) T cells with Th2 phenotypic properties (CD4<sup>+</sup>CXCR4<sup>+</sup>CCR6<sup>-</sup>). Means ± SD; \*p<0.002 compared to Isotype; <sup>‡</sup>p=0.028 compared to IL-6. Pancreatic tissue was stained for **C**) CD3<sup>+</sup> T cells

(brown) by IHC and **D**) quantified at 20× magnification. Means  $\pm$  SD; \*p<0.05 compared to Isotype; <sup>‡</sup>p=0.052 compared to IL-6.

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### Figure 9. IL-6 and PD-L1 blockade increases overall survival of KPC-Brca2 mice

KPC-Brca2 beginning at 5 weeks of age mice were treated with isotype control antibodies or antibodies targeting IL-6 and PD-L1 (200µg/each) until mice were moribund and met pre-specified IACUC-approved early removal criteria. Kaplan-Meier survival curves with log-rank test for significance between isotype control and IL-6/PD-L1 antibodies (p=0.0012).