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CD40 stimulation obviates innate sensors and drives T cell immunity in cancer

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SUMMARY

Cancer immunotherapies are more effective in tumors with robust T cell infiltrates, but mechanisms to convert T cell-devoid tumors with active immunosuppression to those capable of recruiting T cells remain incompletely understood. Here, using genetically engineered mouse models of pancreatic ductal adenocarcinoma (PDA), we demonstrate that a single dose of agonistic CD40 antibody with chemotherapy rendered PDA susceptible to T cell-dependent destruction and potentiated durable remissions. CD40 stimulation caused a clonal expansion of T cells in the tumor, but the addition of chemotherapy optimized myeloid activation and T cell function. Although recent data highlights the requirement for innate sensors in cancer immunity, these canonical pathways – including TLRs, inflammasome, and Type I interferon/STING – played no role in mediating the efficacy of CD40/chemotherapy. Thus, CD40 functions as a non-redundant mechanism to convert the tumor microenvironment immunologically. Our data provide a rationale for a newly initiated clinical trial of CD40/chemotherapy in PDA.

Graphical abstract

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K.T.B. performed and analyzed the experiments, K.T.B. and R.H.V. conceived of and designed the experiments, interpreted data, and wrote the manuscript.



INTRODUCTION

Innate immune cells utilize a number of receptors to detect danger signals liberated when large numbers of host cells die, such as after chemotherapy or radiotherapy in patients with cancer (Green et al., 2009). Dying tumor cells release intracellular components such as high-mobility-group box 1, ATP, and DNA, which are recognized, in turn, by receptors such as Toll-like receptor (TLR) 4 (Apetoh et al., 2007), P2X7 receptor (P2X7R) (Ghiringhelli et al., 2009), and stimulator of interferon genes (STING) (Deng et al., 2014) to regulate immune responses against tumors. Accordingly, a number of innate sensor agonists are being brought forward for investigation in cancer patients (Corrales and Gajewski, 2015; Kaczanowska et al., 2013; Rook et al., 2015).

It is well-known that some chemotherapies can enhance anti-tumor immunity, working most effectively in immunocompetent vs. deficient hosts (Emens and Middleton, 2015; Zitvogel et al., 2008); however, some tumors, such as pancreatic ductal adenocarcinoma (PDA), are notoriously resistant to chemotherapy and despite aggressive treatment, the 5-year survival rate for patients with metastatic PDA is less than 5%. Immunologically, PDA is uncommonly infiltrated by effector T cells and expresses a relatively low burden of nonsynonymous mutations that could serve as neo-epitopes (Alexandrov et al., 2013; Jones et al., 2008; Sausen et al., 2015), consistent with what has been termed an immunologically 'cold' tumor (Sharma and Allison, 2015). Newer combinations of chemotherapy, such as gemcitabine (Gem) and nab-paclitaxel (nP), have shown clinical promise in metastatic PDA (garnering FDA approval in 2013), but objective tumor response rates remain low (23% of patients respond to Gem/nP, compared to 7% with Gem alone) (Von Hoff et al., 2013). Multiple hypotheses have been proposed to explain how nP improves responses against PDA, including SPARC-dependent (Alvarez et al., 2013; Von Hoff et al., 2011) or independent (Neesse et al., 2014) mechanisms of stromal destruction, decreased levels of cytidine deaminase (Frese et al., 2012), and macropinocytosis by Kras-mutant tumor cells

(Commisso et al., 2013). Although paclitaxel may activate macrophages as an LPS mimetic that binds TLR4 (Ding et al., 1993) – which raises the hypothesis of an immune effect from adding nP – progression-free survival is extended by only 1.8 months with Gem/nP compared to Gem alone (Von Hoff et al., 2013) and without durable remissions in this disease.

To investigate immune mechanisms that could convert PDA tumors from T cell-devoid to T cell-replete – as a first step toward establishing immune sensitivity – we used the genetically engineered KPC mouse model of PDA, in which oncogenic Kras^{G12D} and mutant p53^{R172H} are under the control of Cre recombinase specifically expressed in the pancreas. KPC mice develop spontaneous PDA with 100% penetrance and faithful recapitulation of key features of human disease (Hingorani et al., 2005), including a dearth of non-synonymous mutations (similar to other Kras-induced mouse models of cancer (Westcott et al., 2015)) and minimal effector T cell infiltration (Clark et al., 2007). Although CD40 ligation enhances immune activation and maturation of antigen presenting cells (APCs) (Bennett et al., 1998; Ridge et al., 1998; Schoenberger et al., 1998), in tumor-bearing KPC mice, aCD40 alone achieves only transient tumor regressions on the basis of macrophage re-education and not T cell immunity (Beatty et al., 2011). Because α CD40 combined with vaccines drives cytotoxic CD8⁺ T cell responses in the context of cancer (Diehl et al., 1999; French et al., 1999; Sotomayor et al., 1999), we explored aCD40 combined with chemotherapy as an in vivo vaccine (Nowak et al., 2003) against PDA. The inability of α CD40 (with or without Gem) to generate potent T cell mediated regressions of KPC tumors is mitigated upon the depletion of suppressive macrophage populations (Beatty et al., 2015). We hypothesized that adding nP to aCD40/Gem, taking advantage of potential immune stimulating effects of paclitaxel (Ding et al., 1993), might reeducate the suppressive macrophages and promote robust antitumor T cell immunity, bypassing the need for macrophage depletion in this system.

Here, we report that α CD40 and the combination of Gem/nP – but neither α CD40 nor chemotherapy alone – achieves T cell-dependent regression of established tumors in mice, an effect that requires IFN- γ and host CD40. Tumor regression was notably independent of multiple innate sensing pathways that have been classically described as mediating both spontaneous and therapy-induced cancer immunity. These preclinical data provide the mechanistic rationale for a newly initiated clinical trial of Gem/nP/CD40 therapy in patients with PDA (Clinicaltrials.gov, #NCT02588443).

RESULTS

Chemotherapy requires the addition of α CD40 for regression and cure of established PDA in a T cell dependent manner

We harvested a spontaneous PDA tumor from a C57BL/6 KPC mouse and generated a cell line (4662) with mutant Kras and p53 that grew progressively upon subcutaneous implantation in wild-type syngeneic hosts with extensive desmoplastic stroma in the tumor microenvironment (TME) (Lo et al., 2015). Treatment of established 4662 tumors on day 12 with α CD40 and Gem/nP achieved significant regressions 12–14 days later (median regression rate across experiments, 59.7% +/– 26.0%), whereas only rare regressions were observed in mice treated with Gem/nP or α CD40 alone (Figure 1A). Additionally, the

overall tumor growth rate was significantly reduced in Gem/nP/ α CD40 treated mice compared to mice treated with Gem/nP or aCD40 alone (Figure 1B). Similar results were found with a second desmoplastic PDA cell line (G43) also derived from a C57BL/6 KPC mouse (Figure S1). Gem/nP/aCD40 treated mice had significantly enhanced overall survival, with 14.7% (vs. 0%) of mice being cured (Figure 1C). A second dose of Gem/nP 7 days later (day 19), to mimic the weekly dosing schedule in the clinic (Beatty et al., 2013; Von Hoff et al., 2013), neither enhanced nor hindered the rate of regression (Figure S2). Mice that were cured of the primary tumor with Gem/nP/ α CD40 treatment rejected both 4462 and G43 tumor cells when injected 60 days or more later (Figure 1D and data not shown). This effect reflected T cell-mediated memory against PDA, as mice cured with Gem/nP/ α CD40 and then depleted of CD8⁺ T cells after 60 days quickly succumbed to tumor if rechallenged (Figure 1D). Depletion of either CD4⁺ or CD8⁺ T cells, or both, before initial treatment with Gem/nP/ α CD40 also abrogated the response to therapy (Figure 1E). Thus, in contrast to the macrophage-dependent response generated with aCD40 monotherapy, the combination of both Gem/nP and aCD40, but neither alone, effectively mediated T cell-dependent regressions of PDA, reducing overall tumor growth and enabling long-term cures.

Gem/nP/aCD40 therapy skews the PDA microenvironment in favor of effector T cells

Given that T cells mediated tumor regressions prominently on day 23–25, we investigated CD4⁺ and CD8⁺ T cell subsets in the TME at this time point. The prevalence of effector T cells was similar or slightly increased with Gem/nP/ α CD40 compared to Gem/nP or α CD40 alone (Figure 2A), but FoxP3⁺ T regulatory cells (T_{Regs}), comprising nearly 20% of total CD3⁺ T cells in vehicle or Gem/nP treated mice (Figure 2B and data not shown), was significantly reduced after treatment with α CD40, and nearly completely eliminated with the addition of Gem, nP, or both (Figure 2B). As a result, the effector T cell:T_{Reg} ratios were significantly skewed in favor of both CD4⁺ and CD8⁺ effector T cells in the TME after α CD40 (Figure 2C), independently of the addition of Gem and/or nP. The significant reduction in T_{Regs} after α CD40 therapy was observed in both the proportions and in the absolute number of T cell subsets (Figure2B and 2C, and data not shown). CD4⁺ T cell subsets in the TME were significantly altered as early as five days after α CD40, when the proportions of FoxP3⁺ and GATA3⁺ CD4⁺ T cells were significantly reduced in Gem/nP/ α CD40 treated mice, concurrent with an increase in ROR γ t⁺ and Tbet⁺ CD4⁺ cells (Figure 2D).

aCD40 therapy increases the clonal T cell response against PDA

To further investigate the effects of α CD40 on the T cell repertoire, we performed T cell receptor (TCR)- β chain CDR3 region deep sequencing to track unique T cell clones in tumors harvested from mice treated with Gem/nP/ α CD40, α CD40 alone, Gem/nP alone, or vehicle control. To differentiate the effect of each therapy on the TCR repertoire, mice were grouped by α CD40 treatment (Figure 2E–F, top) or Gem/nP treatment (bottom), and analyzed by machine learning using Random Forest classification (RFC), as we have previously reported (Twyman-Saint Victor, et al., 2015). This unbiased analysis approach successfully segregated mice based on α CD40 therapy, regardless of Gem/nP treatment, indicative of the impact of CD40 stimulation (but not chemotherapy) on clonal T cell

responses in the TME. Among all mice that received aCD40, the cumulative proportions of rare and small clones (those found at a frequencies below <0.01%) were significantly increased, and hyperexpanded clones (highly represented clones in the TME) were moderately increased, compared to mice that did not receive aCD40 (2E, top). In comparison, the cumulative frequencies of rare to hyperexpanded clones remained constant when mice were segregated by chemotherapy treatment only, regardless of CD40 treatment (2E, bottom). The moderate increase of hyperexpanded clones in α CD40 treated mice significantly impacted the diversity of the most prevalent clones within the TME, such that the true diversity (measuring the effective number of clones) was increased for the top 10 and 20 clones within the TME, but not for the entire T cell population (Figure 2F, top). Thus the Gini Coefficient (clonality) was significantly increased for the entire response after α CD40 therapy (Figure 2F, top), reflecting the expansion of the most frequent clones in the TME. Again, only exposure to α CD40 and not chemotherapy impacted these diversity and clonal metrics (Figure 2F, bottom). Furthermore, these changes were only observed in the TME itself, using the same machine-learning analysis in the spleen revealed no changes in the clonality or diversity of the T cell repertoire with either Gem/nP or aCD40. Therefore, aCD40 was independently associated with two significant changes in the TCR repertoire specifically within the TME: expansion of certain T cell clones and recruitment of new populations of rare and small clones to the TME.

Functional effector T cells require both Gem/nP and aCD40 treatment

Although α CD40 independently mediated alterations in CD4⁺ and CD8⁺ T cell subsets, the addition of Gem/nP was required for increased functionality of the T cell compartment and control of tumor growth. CD4⁺ T cell production of IFN- γ , TNF- α , and IL-2 was significantly increased in Gem/nP/ α CD40 treated tumors compared to other groups (Figure 2G, left). Moreover, a higher proportion of CD8⁺ T cells produced TNF- α or IFN- γ , or both cytokines, from tumors of mice treated with Gem/nP/ α CD40 compared to Gem/nP or α CD40 alone (Figure 2G, right). Thus, α CD40 significantly reduced the T_{Reg} population and enhanced Th1 and Th17 subsets of CD4⁺ T cells, but the development of functional effector CD4⁺ and CD8⁺ T cells was dependent on the addition of Gem/nP to α CD40.

IFN- γ is required for Gem/nP/aCD40 efficacy

Given the increase in IFN- γ production by both CD4⁺ and CD8⁺ T cells, we investigated the role of IFN- γ in mediating Gem/nP/ α CD40-treatment induced immune responses to PDA. In IFN- γ KO hosts bearing established tumors, response to Gem/nP/ α CD40 therapy at 24 days was fully abrogated (Figure 2H, left). Although vehicle-treated tumors grew somewhat faster in IFN- γ KO mice versus wild-type mice, there was no reduction in tumor growth rate when IFN- γ KO mice were treated with Gem/nP/ α CD40 (Figure 2H, right). IFN- γ is unlikely to be derived from the natural killer cell compartment because depletion with α NK1.1 did not alter tumor responses or growth rates in Gem/nP/ α CD40 treated mice (Figure S3). Additionally, the intratumoral T_{Reg} compartment in IFN- γ KO mice was not significantly reduced after treatment with Gem/nP/ α CD40 as it is in wild-type mice, and consequently the CD8⁺ T cell:T_{Reg} ratio was not skewed in favor of effector T cells (Figure 2I), indicating a failure to generate effector T cells. The potent immune response generated against PDA after

Gem/nP/ α CD40 was therefore dependent on IFN- γ for mediating tumor regressions, and for skewing the tumor microenvironment in favor of effector T cells.

Host CD40 requirement and increased activation of antigen-presenting cells after treatment with Gem/nP/aCD40

To test the mechanism by which CD40-induced immunity is potentiated by Gem/nP, we treated tumor-bearing CD40 knockout (KO) mice with Gem/nP/ α CD40 therapy, and observed no tumor regressions or reduction in overall tumor growth rates (Figure 3A). T_{Reg} reduction and skewing towards effector CD8 T cells at day 24 was also lost in the absence of host CD40 (Figure 3B). Because CD40 KO hosts lack functional germinal center formation for the generation of thymus-dependent B cell responses, we evaluated whether the lack of Gem/nP/ α CD40 efficacy in CD40 KO hosts was due to a defect in the B cell compartment. We measured tumor response rates and growth rates in μ MT KO mice (which lack mature B cells) but found these were similar to those in wild-type mice (Figure 3C). Thus, host expression of CD40 is required for the efficacy of Gem/nP/ α CD40 therapy.

Following treatment with Gem/nP, tumor cell death increased six hours later (Figure S4) (Frese et al., 2012), suggesting potential liberation of tumor antigens *in vivo* prior to α CD40. Three days after chemotherapy administration (24 hours after α CD40), the proportions of activated, MHCII⁺ CD86⁺ CD11b⁺ myeloid cells in the TME were significantly increased in Gem/nP/ α CD40 treated mice compared to other groups, including α CD40 alone (Figure 3D). This increase was also observed in CD11b⁺ F4/80⁺ macrophages and CD11b⁻ CD11c⁺ dendritic cells (DCs) from Gem/nP/ α CD40 treated mice, for which activated populations increased compared to Gem/nP or α CD40 alone (Figure 3A), and was mostly lost by day five (72 hours after α CD40 administration) (data not shown). The proportion of DCs, myeloid cells, and macrophages producing IL-12 in Gem/nP/ α CD40 treated mice was also increased compared to Gem/nP or α CD40 alone (Figure 3E). We observed a concomitant decrease in IL-10 production by CD11b⁺ F4/80⁺ TAMs, CD11b⁺ Ly6C⁺ Ly6G⁺ myeloid derived suppressor cells, and Ly6C^{hi} CD11b⁺ inflammatory macrophages (Figure 3F). Thus Gem/nP/ α CD40 therapy uniquely and significantly enhanced the activation status and function of APCs and myeloid cells in the TME.

Batf3⁺ DCs mediate Gem/nP/aCD40 efficacy

To ascertain the role of APC subsets in mediating Gem/nP/ α CD40 tumor regression, we treated Batf3 KO mice (which lack cross-presenting CD8 α^+ DCs) with Gem/nP/ α CD40, and observed no tumor regressions and a significant diminution in overall tumor growth control (Figure 3G). We also targeted the phagocytic and myeloid cell populations using seven independent depletion methods including clodronate-encapsulated liposomes (Table S1), and although we observed a 30%–50% reduction in the target cell populations in the TME, we were unable to detect any change in treatment efficacy (data not shown and Winograd et al., 2015). Therefore, the effect of Gem/nP/ α CD40 therapy required cross-presentation of tumor antigens by DCs for optimal immune responses against PDA.

Gem/nP/aCD40 therapy drives CD8⁺ T cell-mediated regression of spontaneous PDA

Although 4662 subcutaneous tumors grow with extensive desmoplastic stroma reminiscent of primary PDA (Lo et al., 2015), we also studied Gem/nP/aCD40 treatment against autochthonous tumors arising spontaneously in KPC mice. Mice were enrolled after the diagnosis of a tumor (median volume 103mm³, range 30-400mm³) and treated with Gem/nP on day 0 and day 7, and aCD40 on day 2. Tumor-bearing KPC mice treated with Gem/nP/ α CD40 exhibited a 35.7% total response rate, with tumor regressions in 3/14 mice and stable disease in 2/14 (Figure 4A). In comparison, KPC mice treated with vehicle control, or with the combination of Gem/nP, had no regressions or stabilization of disease, and only 1/14 mice treated with aCD40 alone had stable disease at the 14-day time point after the start of therapy (Figure 4A). The previously reported 30% rate of regressions to aCD40 observed in tumor-bearing KPC mice (Beatty et al., 2011) was not observed here using KPC mice that are fully C57BL/6 backcrossed, although a recent report confirms the macrophagedependency of aCD40 monotherapy in this strain of mice (Long et al., 2016). Moreover, in contrast to the previously reported macrophage-dependent (T cell-independent) regressions in KPC mice treated with CD40 alone (Beatty et al., 2011), here the response rate was completely lost if mice treated with Gem/nP/aCD40 were first depleted of CD8⁺ T cells (Figure 4A), indicating a shift to a T cell-dependent immune response against spontaneous PDA when combining both Gem and nP with aCD40.

CD8⁺ T cell infiltration of the spontaneous PDA TME was significantly increased in KPC mice treated with Gem/nP/ α CD40 as compared to Gem/nP or α CD40 alone (Figure 4C, quantified on right). Additionally, the number of tertiary lymphoid structures (a biomarker of increasingly appreciated immunological importance (Dieu-Nosjean et al., 2014; Lutz et al., 2014)) was significantly increased in spontaneous PDA tumors after Gem/nP/ α CD40 (Figure 4D). The combination of Gem/nP/ α CD40 therapy therefore promotes the development of a robust and orchestrated immune response within the primary tumor site, and allows for CD8⁺ T cell infiltration and destruction of spontaneous KPC tumors, a notoriously difficult site for adaptive immune cells to penetrate.

Gem/nP/aCD40 therapy does not require innate immune sensors for efficacy

Because α CD40 can synergize with TLR agonists (Ahonen et al., 2008) and paclitaxel is an LPS mimetic (Ding et al., 1993), we initially hypothesized that Gem/nP/ α CD40 efficacy would require TLR4 signaling. We were further attracted to this hypothesis because of previous landmark studies reporting a critical role of TLR4 for chemotherapy-induced anti-tumor immunity (Apetoh et al., 2007). However, when TLR4 KO mice were treated with Gem/nP/ α CD40, tumor response rates and growth rates were similar to wild-type mice (Figure 5A). Additionally, robust responses to Gem/nP/ α CD40 were also observed in TRIF KO and MyD88 KO mice, indicating that the downstream mediators of TLR4 (as well as all other TLRs) were not required for therapeutic efficacy (Figure 5B–C). Caspase 11 (Casp 11) can also function as an intracellular LPS receptor (Shi et al., 2014), but Gem/nP/ α CD40 regressed PDA tumors in Casp 11 KO mice the same as wild-type mice (Figure 5D).

Previous reports have shown that ATP released from dying tumor cells stimulates DCs via ATP binding to P2X7R resulting in Casp 1 activation and NLRP3 inflammasome assembly

(Ghiringhelli et al., 2009), but P2X7R KO mice bearing PDA tumors responded similarly to Gem/nP/αCD40 therapy as wild-type hosts (Figure 5E). Additionally, we treated tumorbearing IL-1R KO and Casp 1/11 double KO hosts, and found IL-1 signaling was dispensable for treatment efficacy (data not shown). Therefore, we found no role for the inflammasome pathways in mediating the efficacy of Gem/nP/αCD40 therapy.

Previous studies have shown that MyD88/TLR4/P2X7R pathways are not obligatory for immune responses towards tumors in every setting, but rather, the STING pathway can mediate spontaneous or radiation-induced T cell responses against tumors (Deng et al., 2014; Woo et al., 2014). However, STING mutant (STING Mut) mice, which lack STING function, exhibited tumor response rates and growth kinetics similar to wild-type mice after Gem/nP/aCD40 therapy (Figure 5F). Moreover, Type I IFNs (the downstream target of STING activation) were also dispensable as IFNAR KO hosts responded as well as wildtype hosts to Gem/nP/ α CD40 therapy (Figure 5G), despite the role of Type I IFNs in aCD40/TLR agonist peptide vaccines (Ahonen et al., 2004). To exclude the possibility of cancer cell-autonomous signaling of Type I IFNs (Sistigu et al., 2014), we also blocked IFNAR using anti-IFNAR1 mAb, and found no reduction in the efficacy of Gem/nP/aCD40 therapy (data not shown). Therefore we identified no role for STING or downstream Type I IFNs in mediating responses to Gem/nP/aCD40 therapy. We next investigated IL-12, using IL-12p40 or IL-12p35 KO mice, as well as TNF-a, and found that treatment with Gem/nP/ aCD40 resulted in tumor responses and growth kinetics similar to wild-type mice receiving therapy (Figure 5H and data not shown).

Thus, Gem/nP/ α CD40 treatment is mediated by CD40 and IFN- γ , but independent of 11 other signaling pathways and cytokines, summarized in Table 1. These data illustrate the potency of CD40 stimulation, in combination with Gem/nP, as a non-redundant pathway with the capacity to override the need for classically described innate sensors in mediating activation of anti-tumor immune responses.

Discussion

Although innate immune sensors can play critical roles in spontaneous and therapeutic tumor immunity, here we demonstrate that CD40 stimulation bypasses the need for TLRs, the inflammasome, Type I IFNs, and STING to generate effective priming of adaptive T cell responses against cancer. Using a mutant Kras-driven mouse model of PDA, we observed that treatment with an agonistic α CD40 mAb and chemotherapy alters multiple dimensions of the cancer immunity cycle away from immunosuppression and towards T cell-dependent tumor rejection. The ultimate effect is conversion of an otherwise immunologically 'cold' tumor into one with robust T cell infiltration. Mechanistically, α CD40 and chemotherapy activated myeloid cells and drove T cell function, but α CD40 was required to change T cell profiles in the TME and drive expansion of clonal effector T cell responses. Studies using KO mice showed that host CD40 expression is required for efficacy, as is IFN- γ and cross-presenting DCs. Thus, both gain-of-function and loss-of-function studies highlight the non-redundant role of CD40 activation, in combination with Gem/nP, to obviate the need of innate immune sensing for durable anticancer T cell immunity.

In contrast to immune checkpoint antibodies that unlock pre-existing T cell immunity against cancer, our data support the notion of α CD40 mAb as a complimentary therapeutic strategy in which immune cells are directly activated using an agonistic mAb (rather than blocking mAb) to achieve T cell priming. Expressed by APCs, CD40 uniquely sits proximal in the T cell activation cascade compared to other activation receptors, such as OX40, GITR, or CD137, the ligands of which are upregulated by CD40 activation (Summers deLuca and Gommerman, 2012). To exploit this pathway pharmaceutically, a number of agonistic CD40 antibodies are being evaluated in cancer clinical trials (Melero et al., 2013; Vonderheide and Glennie, 2013). Our group has shown that one such CD40 mAb (CP-870,893) results in modest rates of objective tumor regression as a single agent in patients with melanoma (Bajor et al., 2014; Vonderheide et al., 2007) in the absence of autoimmune-like events associated with aPD-1 or aCTLA-4 therapy. Nevertheless, studies from tumor-bearing mice predict that aCD40 alone in the absence of a "vaccine" to deliver tumor antigen will be an inefficient therapeutic approach. Indeed, T cell-mediated tumor regressions with aCD40 alone in mice have largely been reported only in immunogenic tumors such as those expressing viral antigens (van Mierlo et al., 2002).

We therefore examined the therapeutic prospect of α CD40 as an immune combination partner in our PDA models with a new standard-of-care Gem/nP chemotherapy. Although the addition of Gem to aCD40 was found to enable T cell immunity against murine mesothelioma (Nowak et al., 2003), in our model of PDA, Gem/aCD40 (without nP) mediates potent T cell immunity against subcutaneous tumors, but not in spontaneous KPC tumors for which the T cell response is restrained by macrophages (Beatty et al., 2011; Beatty et al., 2015). Accordingly, Gem/aCD40 therapy resulted in modest tumor regression rates in patients with metastatic PDA, but tumors lacked T cell infiltrates, and all patients eventually progressed (Beatty et al., 2011). Here, using the chemotherapy doublet of Gem/nP, we observed clear evidence of T cell-mediated regression in both subcutaneous and spontaneous KPC tumors, suggesting an immunological benefit of Gem/nP compared to Gem alone. Probing the immunological mechanism underlying Gem/nP/CD40 efficacy, we found that chemotherapy and α CD40 therapy shifted the myeloid compartment toward an M1 bias, and the T cell subsets toward a Th1 bias, in terms of both phenotype and function, with a near complete collapse of the intratumoral TReg compartment. Importantly, based on TCR deep sequencing of intratumoral T cells, treatment with aCD40 was independently associated with expansion of the top clones within TCR repertoire, as well as the recruitment of new clones to the TME.

Taken together, our findings support a mechanistic model of tumor immunity in which the addition of both Gem and nP converts the effect of α CD40 therapy from macrophage-dependency to T cell-dependency. The combination chemotherapy fuels tumor antigen release that cooperates with CD40-mediated DC activation and drives T cell priming. nP, but not Gem, increased tumor cell death shortly after administration so that α CD40 given two days later optimally impacts antigen-loaded DCs. Moreover, efficacy of Gem/nP/ α CD40 was lost in Batf3 KO mice, which lack DCs most capable of antigen cross-presentation. Thus, insufficient APC activation and antigen presentation – an important immune deficiency in cancer – may be uniquely addressed via α CD40 therapy.

Given the immune benefit from the addition of Gem/nP, it is interesting that classical innate immune sensing – as evaluated *in vivo* both genetically and pharmacologically – played no role in mediating T cell regression triggered by Gem/nP/αCD40. Eleven such pathways – including MyD88, P2X7R, and IFNAR – were tested, but none was found to be required. In certain previous studies, chemotherapy alone induces immunogenic tumor cell death dependent on host MyD88/TLR4 signaling (Apetoh et al., 2007). In other experimental models, response to chemotherapy is independent of the adaptive immune system, particularly in spontaneous mouse tumor models (Ciampricotti et al., 2012), and may require additional modifiers of the TME to trigger T cell responses, e.g. inhibition of CSF-1R (DeNardo et al., 2011; Zhu et al., 2014) or BTK (Masso-Valles et al., 2015).

Our approach with α CD40 is therapeutically and mechanistically distinct from other strategies to enhance T cell immunity against PDA, offering the potential for further synergistic combinations. For example, FAP⁺ stromal cells in PDA regulate T cell infiltration to PDA via CXCL12/CXCR4 (Feig et al., 2013), but FAP⁺ stromal cells in the KPC model are CD40-negative, and FAP⁺ cell depletion (or CXCR4 inhibition) does not negatively impact T^{Regs} in the way aCD40/chemotherapy does in the same KPC model. Vaccination with recombinant antigen-expressing Listeria is another powerful method to generate anti-PDA T cells (Keenan et al., 2014) but appears to rely on STING activation (Jin et al., 2013; Woodward et al., 2010), unlike aCD40. Other treatments that can mediate T cell responses against PDA include GVAX vaccination (Le et al., 2015; Soares et al., 2015), adoptive transfer of antigen-receptor engineered T cells (Stromnes et al., 2015), and CSF-1R inhibition (Zhu et al., 2014). Although antibody blockade of PD-1 (or PD-L1) with or without aCTLA-4 is largely ineffective in treating PDA in mice or patients (Brahmer et al., 2012; Herbst et al., 2014; Twyman-Saint Victor et al., 2015; Winograd et al., 2015; Zhu et al., 2014), PD-1 blockade in mice synergizes with certain T cell therapies in PDA (Feig et al., 2013; Le et al., 2015; Soares et al., 2015; Zhu et al., 2014). Indeed, we have shown that the addition of checkpoint blockade to Gem/nP/aCD40 in tumor-bearing mice enhances survival in both implantable and spontaneous PDA models (Winograd et al., 2015), and here we show the mechanism by which the PDA TME is rendered sensitive to PD-1 and CTLA-4 antibodies used in that study. Taken together, these reports highlight multiple immune vulnerabilities of PDA that can be targeted in a non-redundant fashion in combination with aCD40 in clinical trials (Melero et al., 2013).

Although the T cells generated by Gem/nP/ α CD40 mediate tumor regressions and long-term protection, the precise antigens targeted by this response remain unknown. The minimal expression of non-synonymous mutations in our KPC model and the lack of predicted neo-epitopes able to bind MHC class I (n=0–5 predicted neo-epitopes per tumor; unpublished data) suggests the target peptide-MHC tumor repertoire is mechanistically distinct from that underlying responsiveness to checkpoint blockade. Human PDA also exhibits a scarcity of non-synonymous mutations such that the burden of neo-epitopes may be relatively low compared to carcinogen-induced tumors such as lung carcinoma or melanoma (Alexandrov et al., 2013; Gubin and Schreiber, 2015; Jones et al., 2008; Sausen et al., 2015). Although peptides derived from mutated Kras can potentially function as tumor-specific antigens (Tran et al., 2015), vaccination against mutated Kras is unable to slow growth of established PDA tumors (Keenan et al., 2014). It is possible that T cells generated after Gem/nP/ α CD40

treatment are specific for self-antigens, but we did not observe autoimmunity or related toxicities in our experiments, suggesting that these potential antigens are not strongly expressed on essential tissues. Given the shared protection between two independent KPC-derived PDA cell lines, our findings justify a reconsideration of self-antigens – as well as "abnormal self-antigens" not derived on the basis of non-synonymous mutations (Cobbold et al., 2013; Ryan et al., 2010) – as potential tumor rejection antigens.

In summary, our findings demonstrate the powerful ability of a single dose of α CD40 to alter T cells in the TME, expand clonal T cell populations, and convert the TME in pancreatic cancer to a site replete with infiltrating T cells. In combination with a novel chemotherapy doublet, α CD40 treatment bypasses innate immune sensors to generate functional APCs and T cells, culminating in durable responses with curative potential, even in a highly immunosuppressive TME. With the goal of rapidly translating these observations to patients, a newly opened clinical trial (Clinicaltrials.gov, #NCT02588443) is evaluating administration of Gem/nP and α CD40 before and after surgery in patients presenting with resectable PDA.

EXPERIMENTAL PROCEDURES

Mice

Kras^{LSL-G12D/+},Trp53^{LSL-R172H/+}, Pdx1-Cre (KPC) mice have been previously described (Hingorani et al., 2005), and were bred and maintained in the specific pathogen-free facility at the University of Pennsylvania. The genetic background of the C57BL/6 KPC mice was assessed at the DartMouse[™] Speed Congenic Core Facility at the Geisel School of Medicine at Dartmouth College, as described in Supplemental Information. STING Mut (*Tmem173^{gt}/J*) (Sauer et al., 2011) were kindly provided by Dr. Susan Ross (Perelman School of Medicine, University of Pennsylvania). All wild-type C57BL/6 and other KO mice (Supplemental Information) were purchased from The Jackson Laboratory and/or bred at the University of Pennsylvania. Most experiments with wild-type C57Bl/6 mice were performed in female mice, but tumor growth responses were confirmed in male mice. Experiments in KO and KPC mice were performed with mixed gender mice distributed across treatment groups. Animal protocols were reviewed and approved by the Institute of Animal Care and Use Committee at the University of Pennsylvania.

Cell Lines and in vivo growth

The mouse pancreatic cancer cell line 4662 was previously described (Lo et al., 2015). PDA cells were used in experiments after 3–5 passages *in vitro;* C57Bl/6 mice received 2.5×10^5 PDA cells subcutaneously only if tumor cell viability was >94%. Cell lines were tested by using the Infectious Microbe PCR Amplification Test (IMPACT) and authenticated by the Research Animal Diagnostic Laboratory (RADIL) at the University of Missouri. Tumors were measured thrice weekly by calipers, and the volume was calculated by (LxW²)/2, where L is the longest diameter and W is the perpendicular diameter. Mice were designated as responders if tumors had regressed 12–14 days after the initiation of treatment.

Drug preparation

Gemcitabine (Gem; Hospira) pharmaceutical grade suspension at 38mg/mL 2'-deoxy-2',2'difluorocytidine was diluted to 12mg/mL in PBS and administered at 120mg/kg via intraperitoneal (i.p.) injection (Beatty et al., 2011). Nab-paclitaxel (nP; Abraxane, Celgene) pharmaceutical grade powder was resuspended at 12mg/mL in PBS and administered at 120mg/kg i.p. (Frese et al., 2012) or equivalent molar dose of human albumin (huAlb) (Sigma). Gem/nP or huAlb was injected on day 12 after tumor injection in subcutaneous PDA experiments, and on days 0 and 7 in KPC mice. Gem and nP were purchased through the Hospital of the University of Pennsylvania Pharmacy.

Monoclonal antibodies

Mice received 100 µg of either agonist CD40 rat anti-mouse IgG2a mAb (clone FGK45, endotoxin-free), or the isotype control IgG2a mAb (clone 2A3) (Beatty et al., 2011) on day 14 after 4662 injection, or day 2 in KPC mice. CD4⁺ or CD8⁺ T cells were depleted with 200 µg each of clone GK1.5 or clone 2.43, respectively, injected i.p. on day 10 and repeated every 4 days, or IgG2b isotype control (clone LTF-2). CD4⁺ and CD8⁺ T cell depletion was confirmed by staining peripheral blood (data not shown). KPC mice received CD8 depleting antibody starting on day –1 and repeated every 4 days until day 14. All antibodies were purchased from BioXCell.

Tumor regression studies in KPC mice

KPC mice were monitored for spontaneous tumors by ultrasonography every 1–2 weeks using the Vevo 2100 Imaging System with 55MHz MicroScan Transducer from Visual Sonics. Mice with tumors measuring at least 30mm³ were enrolled within 24 hours of baseline imaging using blocked randomization to assign treatment group. Mice were designated as responders if disease was stable (progression <20% compared to baseline) or if tumors regressed 14 days after initiation of treatment.

Preparation of tissue samples from mice

Mice were euthanized either on day 15, 19, 24, or 26 after 4662 injection, and tumors, draining lymph nodes, and spleens were harvested, as indicated. Tumors were minced and incubated for 45 minutes in 1mg/mL collagenase V in DMEM at 37°C. Tumors, spleen, and lymph nodes were mechanically dissociated and passed through a 70 μ M cell strainer, spleens were incubated in ACK lysis buffer (BioWhittaker), and then tissues were used for flow cytometric analysis as single cell suspensions. Cells were counted using the Beckman Coulter Counter Z2.

Flow Cytometry

Cell surface molecules were analyzed by incubating single cell suspensions of tissues with primary fluorochrome-labeled antibodies at 4°C for 30 minutes in PBS with 0.5% BSA and 2mM EDTA. For cytokine production by T cell subsets, samples were incubated for 4–5 hours at 37°C with PMA/ionomycin (Sigma) and Brefeldin A (Sigma). Intracellular staining was done using the fixation/permeabilization kit from eBiosciences. For cytokine production by APCs and myeloid cells, samples were incubated for 4–5 hours with Brefeldin A and

Golgistop (BD Biosciences), with 1µg/mL LPS (Sigma). Antibodies used in flow analysis are described in Supplemental Information. Flow cytometric analysis was performed on a FACSCanto or LSR II flow cytometer (BD Biosciences). Collected data were analyzed using FlowJo software (Treestar).

Immunohistochemistry

Tumors were embedded in OCT and then sectioned in 5um slices, fixed in acetone, and stained using a Bond Max automated staining system (Leica Microsystems), with the Bond Intense R staining kit (Leica Microsystems), using CD8 primary antibody (clone 53-6.7, Abcam). H&E stains were performed according to manufacturer's directions (Sigma). The histopathological scoring is detailed in Supplemental Information.

TCR deep sequencing and analysis

High-throughput next-generation sequencing of the TCR- β CDR3 region was performed by Adaptive Biotechnologies using the ImmunoSeq platform (Supplemental Information). Analysis of TCR- β repertoire was performed using the tcR R package (Nazarov et al., 2015). Random Forest machine learning for classification predictions was performed using the randomForestSRC R package (Ishwaran and Kogalur, 2010) as previously described (Twyman-Saint Victor et al., 2015).

Statistical Analyses

Significance of overall survival was determined using Kaplan-Meier survival curve with Log-rank analysis. All other comparisons were performed using one- or two-way ANOVA with Tukey's HSD post-test, or Mann-Whitney T test, as indicated. All statistical analyses were performed with Graphpad Prism 6 (GraphPad). Standard deviation (SD) or Standard Error of the Mean (SEM) shown as indicated by error bars. P < 0.05 was considered statistically significant, * indicates P < 0.05, ** P < 0.01, *** P < 0.001, and *** P < 0.0001, *ns* (or lack of indicated P value) denotes not significant (P > 0.05).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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- CD40 stimulation converts T cell-deficient tumors to immunologically replete sites
- CD40 mediates clonal T cell expansion with decreased regulatory T cells
- Functional adaptive immune responses require both CD40 stimulation and chemotherapy
- Converted tumors undergo durable responses independently of innate immune sensors

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Figure 1. Gem/nP/aCD40 drives T cell-dependent regressions of PDA

Mice were injected with PDA 4662 cells subcutaneously and after 12 days of growth, tumors were treated with Gem/nP followed by aCD40 2 days later.

(A) Left, change in tumor volume on day 24 compared to start of treatment (day 12), representative of 7 independent experiments. Right, the total proportion of regressors/ experiment, from 13 individual experiments, total number of mice/group shown below.(B) Tumor growth kinetics for mice from (A).

(C) Survival curve for mice treated as described in (A), from 2 combined experiments.

(D) Survival after second tumor injection >60 days after primary tumor injection. Some mice received α CD8, representative of 2 independent experiments.

(E) Mice were treated as described in (A), and with α CD4 and/or α CD8. Left, change in tumor growth compared to baseline, right, tumor growth kinetics. Data are representative of 3 independent experiments.

Each experiment had 4–10 mice/group, each bar represents a single mouse and each symbol represents a group, horizontal line and error bars indicate mean ± SEM. Statistical analyses by one-way ANOVA (A), two-way ANOVA with Tukey's HSD post-test (B, E), or Log-rank test (C, D). See also Figure S1, S2.



Figure 2. Gem/nP/aCD40 therapy alters T cell subsets, repertoire, and function in PDA tumors in an IFN- γ dependent manner

(A–D) Mice were treated as described in Figure 1A, and tumors were harvested 24 days (A–C) or 19 days (D) after tumor injection (12 and 7 days after initiation of treatment,

respectively) and analyzed by flow cytometry with regard to the proportion (A, B, D) of the indicated subsets or the ratios of the absolute number of cells/gram of tumor (C) among live, $CD45^+$ CD3⁺ cells.

(E–F) Tumors were harvested on day 24 and analyzed by TCR deep sequencing. Mice are grouped based on receiving CD40 (top) or Gem/nP (bottom), and the cumulative frequencies

of Rare (representing $<10^{-5}$ total clones), Small (10^{-5} to $<10^{-4}$), Medium (10^{-4} to $<10^{-3}$), Large $(10^{-3} \text{ to } < 10^{-2})$, or Hyperexpanded $(10^{-2} \text{ to } 1)$ clones within the total repertoire are indicated (E), or the repertoire diversity ('true diversity,' indicating effective number of clones) for the top 10, top 20, or entire population (far left to middle right), or the Gini Coefficient (0 indicating polyclonal, 1 indicating monoclonal) on far right (F). (G) Tumors were harvested at day 24 and analyzed by flow cytometry with regard to the indicated parameters among CD4⁺ (left) or CD8⁺ (right) live, CD45⁺ CD3⁺ cells. (H, I) IFN-7 KO mice were treated as described in Figure 1A. (H) Change in tumor volume on day 12 (left) with growth kinetics (right). (I) Tumors were analyzed on day 24 by flow cytometry with regard to the indicated subsets or ratios among live, CD45⁺ CD3⁺ cells. Each symbol represents an individual mouse, horizontal lines indicate mean ± SD (A-G, I) except for (H) where each bar represents a single mouse and each symbol represents a group with mean \pm SEM. Data representative of 3–5 independent experiments with 4–6 mice/ group, except TCR deep sequencing data, 1 experiment with 8-9 mice/group. Statistical analysis was performed by one-way ANOVA (A-D, G, I), Mann-Whitney T test (E, F) or two-way ANOVA (H) with Tukey's HSD post-test. See also Figure S3.



Figure 3. Gem/nP/aCD40 therapy requires host CD40, activates antigen-presenting cells and requires Batf3⁺ dendritic cells for efficacy

Mice were treated as described in Figure 1A.

(A, B) CD40 KO mice. (A) Left, change in tumor volume on day 24 versus day 12 (start of therapy). Right, tumor growth kinetics. (B) Tumors were analyzed on day 24 with regard to the proportions of indicated cells and ratios among live, CD45⁺ CD3⁺ cells.

(C) μ MT KO mice. Left, change in tumor volume on day 24 compared to day 12. Right, tumor growth kinetics.

(D–F) Mice were treated as described in Figure 1A, and tumors were harvested on day 15 (24 hours after receiving CD40) and analyzed by flow cytometry with regards to the proportions of indicated subsets among live, CD45⁺ CD3⁻ cells. CD11c⁺ cells are also CD11b⁻ F4/80⁻.

(G) Batf3 KO mice treated as in Figure 1A. Left, the change in tumor volume on day 24 versus day 12, right, tumor growth kinetics.

Each bar represents an individual mouse, symbols indicate groups, and horizontal lines indicate mean \pm SD (A, C, G) or each symbol represents an individual mouse, with mean \pm SEM (B, D–F), data representative of 2–5 independent experiments with 3–10 mice/group. Statistical analysis by one-way ANOVA (A–C) or two-way ANOVA (D) with Tukey's HSD post-test. See also Figure S4 and Table S1.

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Figure 4. Spontaneous tumors in KPC mice respond to Gem/nP/aCD40 therapy in T cell-dependent fashion

KPC mice diagnosed with established tumors received Gem/nP on day 0 and day 7, and α CD40 was given on day 2. Some mice (as indicated) also received α CD8 depletion for the duration of enrollment. (A) The change in tumor volume on day 14 compared to initial tumor volume at diagnosis, responders calculated in table below.

(B) Tumor growth curves for indicated groups, responders indicated in red.

(C) Representative histological samples from (A) at day 14, stained for CD8, shown on left at two magnifications, quantification of global CD8 staining in tumors on right. Scale bar indicates $200 \ \mu m$ (top) or $300 \ \mu m$ (bottom).

(D) Representative H&E samples of tumors from (A) at day 14 shown on left, quantification of tertiary lymphoid structures (TLS) in entire tumor section on right. Top, $2\times$, Arrowheads point to TLS, asterisk indicated tumor-associated lymph node, and outline indicates field below ($20\times$). Scale bar indicates 1000 µm (top) or 100 µm (bottom).

Each bar, line, or symbol represents an individual mouse, horizontal lines indicate mean \pm SD. Statistical analysis by one-way ANOVA with Tukey's HSD post-test (C, D), and Fisher's Exact Test (A).

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Figure 5. Gem/nP/aCD40 therapy bypasses innate immune sensors for treatment efficacy Mice were treated as outlined in Figure 1A, and for each panel: Left, change in tumor volume on day 24 compared to day 12 (start of therapy), right, tumor growth kinetics. (A) TLR4 KO (B) MyD88 KO (C) TRIF KO (D) Casp 11 KO (E) P2X7R KO (F) STING Mut (G) IFNAR KO and (H) IL-12p40 KO.

Each bar represents a single mouse, each symbol represents a group with error bars indicating mean \pm SEM, data shown representative of 2–5 independent experiments for each KO strain with 4–10 mice per group. Statistical analysis was by two-way ANOVA with Tukey's HSD post-test.

Table 1

Gem/nP/ $\alpha CD40$ is dependent only on host CD40 and IFN- $\gamma,$ T cells, and Batf3^+ DCs

Target	Response to Gem/nP/aCD40
Signaling Molecules	
CD40	No
MyD88	Yes
TLR4	Yes
TRIF	Yes
TLR3	Yes
Caspase 1	Yes
Caspase 11	Yes
STING	Yes
P2X7R	Yes
Cytokines	
IFN-γ	No
IFN- α/β	Yes
IL-1	Yes
IL-2	Yes
TNF-a	Yes
Adaptive Immune Cells	
CD4 ⁺ T cells	No
CD8 ⁺ T cells	No
B Cells	Yes
Innate Immune Cells	
Batf3 ⁺ DCs	No
NK Cells	Yes