

Radiotherapy Combined with Novel STING-Targeting Oligonucleotides Results in Regression of Established Tumors

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Abstract

Cytotoxic therapies prime adaptive immune responses to cancer by stimulating the release of tumor-associated antigens. However, the tumor microenvironment into which these antigens are released is typically immunosuppressed, blunting the ability to initiate immune responses. Recently, activation of the DNA sensor molecule STING by cyclic dinucleotides was shown to stimulate infection-related inflammatory pathways in tumors. In this study, we report that the inflammatory pathways activated by STING ligands generate a powerful adjuvant activity for enhancing adaptive immune responses to tumor antigens released by radiotherapy. In a murine model of

pancreatic cancer, we showed that combining CT-guided radiotherapy with a novel ligand of murine and human STING could synergize to control local and distant tumors. Mechanistic investigations revealed T-cell-independent and TNF α -dependent hemorrhagic necrosis at early times, followed by later CD8 T-cell-dependent control of residual disease. Clinically, STING was found to be expressed extensively in human pancreatic tumor and stromal cells. Our findings suggest that this novel STING ligand could offer a potent adjuvant for leveraging radiotherapeutic management of pancreatic cancer. *Cancer Res*; 76(1); 50–61. ©2015 AACR.

Introduction

Radiation therapy (RT) is a potential means to kill cancer cells to release tumor antigens and endogenous immune adjuvants for immune priming (1, 2), and also to identify the tumor by initiating local inflammation to direct trafficking and function of effector T cells at the treatment site (3–5). In murine models, RT depends in part on T cells for tumor control (6–9). RT synergizes well with T-cell-targeted immunotherapies in murine models (8, 10–14), in early clinical studies (15) and anecdotal reports (16–18). However, the tissue damage caused by RT also initiates a well-characterized wound repair response at the treatment site (19). This wound repair process is driven by intratumoral macrophages that limit inflammation and immune-mediated tissue destruction (20–22). Thus, depletion of tumor macrophages (23, 24) or preventing macrophage polarization to wound-healing phenotypes (25) significantly enhances the efficacy of RT.

We are investigating therapeutic interventions that prevent the postradiation transition to an immune suppressive environment and therefore have the potential to enhance immune-mediated tumor control following RT. Infectious agents are known to prevent wound healing (26) and both infectious agents and immunologic adjuvants have shown synergy with RT (27–30). STING (stimulator of interferon genes) is a cytosolic sensor of microbial infection (reviewed in ref. 31). Importantly, hyperactivity of STING to host DNA can result in autoimmunity (32) and activating mutations in STING can result in immunopathology (33). Recent data demonstrate that host STING expression is critical for spontaneous rejection of immunogenic or partially allogeneic tumors via effects on tumor macrophages (34) and radiation-mediated cure of immunogenic tumors is dependent on host STING (35). We hypothesized that ligands that activate STING will initiate inflammatory mechanisms within the tumor environment to improve the efficacy of RT in unresponsive tumors.

To test our hypothesis, we utilized synthetic CDN derivatives that were designed to have increased activity compared with natural STING ligands produced by bacteria or by host cell cyclic guanosine monophosphate-adenosine monophosphate synthase (cGAS). Since CDNs are susceptible to degradation by phosphodiesterases, dithio analogues in which the nonbridging oxygen atoms in the internucleotide phosphate bridge were replaced with sulfur atoms during synthesis. Since the phosphate bridge constitutes a chiral center, we tested purified dithio diastereomers and found that the R_pR_p molecule had increased immunostimulatory properties *in vitro*, and conferred increased therapeutic antitumor efficacy *in vivo*, as compared with the R_pS_p dithio analogue or the parent CDN compound (Fu and colleagues, manuscript submitted). When the R_pR_p dithio modification was combined with compounds that contained the same 2'-5', 3'-5' noncanonical (or

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mixed) linkage phosphate bridge as 2'-3' cGAMP, we found that these R_p,R_p dithio 2'-3' CDN molecules activated all five human STING alleles (36) expressed in cell lines, activated human PBMCs including from donors homozygous for the refractory allele, and had potent therapeutic efficacy in tumor-bearing mice (37).

We hypothesize that the potent inflammatory response elicited by STING ligation will prevent initiation of the wound-healing process and will enhance adaptive immune responses to antigens released by RT. To test this hypothesis, we use a murine pancreatic cancer model that is poorly responsive to RT and following treatment generates a myeloid response that suppresses adaptive immunity (7, 25, 38). CT-guided radiation was used to target tumors with minimal dose to normal tissues, and combined with local injection of a novel human and murine STING agonist. We demonstrate that RT and STING ligation synergize to generate systemic T-cell responses that control distant disease. We demonstrate that early tumor control by STING ligation is T-cell independent but occurs due to TNF α -mediated hemorrhagic necrosis, while late control of recurrence is CD8 T-cell dependent. These data demonstrate that application of this novel STING ligand can convert radiation-mediated cell death into an endogenous vaccine to enhance adaptive immune-mediated local and distant tumor control.

Materials and Methods

Animals and cell lines

The Panc02 murine pancreatic adenocarcinoma cell line (39) was kindly provided in 2009 by Dr. S. Woo (Mount Sinai School of Medicine, New York, NY). The 3LL lung adenocarcinoma cell line (40) was obtained in 2009 from the ATCC. The SCCVII squamous cell carcinoma cell line was kindly provided in 2014 by Dr. W. Lee (Duke University Medical Center, Durham, NC). Species identity checks on these murine cell lines were performed using murine-specific MHC antibodies, and were tested for contamination within the past 6 months using a Mycoplasma Detection Kit (SouthernBiotech). Six to 8-week-old C57BL/6 (Panc02, 3LL), C3H (SCCVII) or FVB (MMTV-PyMT) mice were obtained from Charles River Laboratories for use in these experiments. STING^{-/-} (Goldenticket Stock# 017537) and Rag1^{-/-} (Stock # 002216) mice were obtained from The Jackson Laboratory. Transgenic MMTV-PyMT mice (41) were kindly provided by Dr. E. Akporiaye (Sidra Medical and Research Center, Doha, Qatar). Mice expressing Cre under the PDX promoter (Stock #014647) and floxed Kras^(G12D) (Stock # 008179) were obtained from The Jackson Laboratory and mice with floxed Trp53^(R172H) (Stock # 01XM2) were obtained from the NCI Fredrick Mouse Repository. Mice were cross-bred to generate Pdx-Cre^{+/-} Kras^{(G12D)+/-} Trp53^{(R172H)+/-} mice that spontaneously generate pancreatic tumors (42). Survival experiments were performed with 6 to 8 mice per experimental group, and mechanistic experiments with 4 to 6 mice per group. Animal protocols were approved by the Earle A. Chiles Research Institute Institutional Animal Care and Use Committee (Animal Welfare Assurance No. A3913-01).

Antibodies and reagents

Depleting anti-CD8 antibody (YTS 169.4 – BioXCell) was given i.p. 250 μ g one day before treatment and again 1-week later. Blocking anti-TNF α antibody (XT3.11 – BioXCell) was given i.p. 500 μ g 6 hours before treatment and again 48 hours later. Fluorescently-conjugated antibodies CD11b-AF700, Gr1-PE-Cy7, IA

(MHC class II)-APC-Cy7, CD4-FITC (HIS51), CD8-PerCP-Cy5.5 (53.6.7), IFN γ -APC(XMG1.2) were purchased from Ebioscience. Western blotting antibodies used include arginase I (BD Biosciences), iNOS (Cayman Chemical Corporation), GAPDH, anti-mouse-HRP, and anti-rabbit-HRP (all Cell Signaling Technology). Immunohistology antibody to F4/80 was purchased from ADB Serotec; CD3, CD31 from Spring Bio; Cy3 conjugated Smooth Muscle Actin from Sigma; CD45 (30-F11) from Ebioscience; and unconjugated anti-STING from Cell Signaling Technology.

Preparation of synthetic STING ligands

Modified CDN derivative molecules were synthesized according to modifications of the "one-pot" Gaffney procedure, described previously (37, 43). Synthesis of CDN molecules utilized phosphoramidite linear coupling and H-phosphonate cyclization reactions. Synthesis of dithio CDNs was accomplished by sulfurization reactions to replace the non-bridging oxygen atoms in the internucleotide phosphate bridge with sulfur atoms. The phosphorus III intermediates generated upon formation of the linear dimer (phosphite triester stage) and cyclic dinucleotide (H-phosphonate diester stage) were sulfurized by treatment with 3-((N,N-dimethylaminomethylidene)amino)-3H-1,2,4-dithiazole-5-thione (DDTT) and 3-H-1,2-benzodithiol-3-one, respectively. The crude reaction mixture obtained after the second sulfurization was chromatographed on silica gel to generate a mixture of the RR- and RS-diastereomers of fully protected ML S2 CDA. Benzoyl and cyanoethyl deprotection using methanol and concentrated aqueous ammonia generated bis-TBS-ML-S2 CDA as a mixture of RR- and RS-diastereomers, which were separated by C-18 prep HPLC. The purified bis-TBS-ML RR-S2 CDA was deprotected with TEA-3HF, neutralized with 1 mol/L triethylammonium bicarbonate and desalted on a C18 SepPak to give ML RR-S2 CDA as the bis-triethylammonium salt in >95% purity. CDN molecules were characterized by high-resolution Fourier transform ion cyclotron resonance mass spectroscopy (FT-ICR) to confirm the expected elemental formula, both ¹H NMR and ³¹P NMR, and by ¹H-¹H COSY (correlation NMR spectroscopy) in combination with a ¹H-³¹P HMBC (heteronuclear multiple-bond correlation spectroscopy) to confirm the regiochemistry of the phosphodiester linkages. Before use in experiments, all synthetic CDN preparations were verified by LAL assay to be endotoxin free (<1 EU/mg).

RT of tumors

Tumors were inoculated in the right flank at a dose of 2×10^5 Panc02, 5×10^5 3LL, and 5×10^5 SCCVII. MMTV-PyMT tumors were established in naïve FVB mammary glands using the published method (44). Briefly, tumors were harvested from day 100 MMTV-PyMT⁺ mice and dissected into approximately 2 mm fragments followed by agitation in 1 mg/mL collagenase in PBS for 1 hour at room temperature. The digest was filtered through 100 μ m nylon mesh to remove macroscopic debris and 1×10^6 cells were injected into the mammary fat pad in a 1:1 mix with Matrigel (BD Biosciences). Ten to 14 days after tumor challenge, mice were randomized to receive treatment with CT-guided RT using a Small Animal Radiation Research Platform (SARRP, XStrahl, Gulmay Medical) to deliver 10 Gy to an isocenter in the tumor, with beam angles designed to minimize dose to normal tissues. Dosimetry was performed using SLICER software with SARRP-specific add-ons (XStrahl). Mice were additionally randomized to receive concurrent intratumoral injection of doses of

RR-CDG in 25 μ L of PBS or PBS vehicle alone, with an additional intratumoral injection 24 hours later.

Tumor-associated antigen-specific responses

Control mice and mice bearing Panc02 expressing the model antigen SIY (Panc02-SIY, kindly provided by Dr. R. Weishelbaum, University of Chicago, Chicago, IL) were randomized to receive treatment as described above. Spleens were harvested 7 days following treatment and cell suspensions were stimulated with 2 μ mol/L of (SIYRYGL) or DMSO vehicle in the presence of brefeldin A for 5 hours at 37°C. Stimulated cells were washed and stained with CD4-FITC and CD8-PerCP Cy5.5, then fixed and permeabilized using a BD Cytofix/Cytoperm Plus Kit (BD Biosciences) and frozen at -80° C. For analysis cells were thawed and intracellularly stained with IFN γ -APC. Cells were washed and analyzed on a BD LSRII. Flow cytometer and the data were interrogated using BD FACSDiva (BD Biosciences) and FlowJo (Tree Star).

Immunohistochemistry

For immunohistology, murine tumors were fixed in Z7 zinc-based fixative (35) overnight. Tissue was then processed for paraffin tissue sections. Five-micrometer sections were cut and mounted for analysis. Tissue sections were boiled in EDTA buffer as appropriate for antigen retrieval. Primary antibody binding was visualized with AlexaFluor 488, AlexaFluor 568, or AlexaFluor 647 conjugated secondary antibodies (Molecular Probes) and mounted with DAPI (Invitrogen) to stain nuclear material. Routinely processed paraffin-embedded tumor tissue and adjacent normal tissue from surgical specimens of patients with confirmed pathologic diagnosis of adenocarcinoma were used to prepare 5- μ m sections of sections for staining. Sections were deparaffinized, rehydrated then deprotected before staining for STING. Antibody binding was detected using HRP-conjugated secondary antibody and DAB as the enzyme substrate; slides were counterstained with hematoxylin, dehydrated and mounted for analysis. Images were acquired using a Zeiss Axio observer Z1 with attached Nuance Multispectral Image camera and software (Perkin Elmer) or a Leica SCN400 whole slide scanner. All images displayed in the manuscript are representative of the entire tumor and their respective experimental cohort.

Cytokine bead assay

Detection of cytokines from murine peripheral blood serum, tumor homogenates, and flow sorted tumor-associated macrophages was performed using a murine multiplex bead assay as previously described (45). To prepare tumor homogenates, tumors were harvested on ice and homogenized in 4.5 μ L PBS containing HALT protease inhibitor per mg tissue. The cell debris was removed by centrifugation at 14,000 g for 15 minutes at 4°C, and supernatants were stored in aliquots at -80° C until used. Cytokine levels in the supernatants were detected using a murine multiplex bead assay (Life Technologies) and read on a Luminex 100 array reader. Cytokine concentrations for replicates of each tumor sample were calculated according to a standard curve.

Flow sorting of tumor macrophages

The tumor was dissected into approximately 2 mm fragments followed by agitation in 1 mg/mL collagenase (Invitrogen), 100 μ g/mL hyaluronidase (Sigma), and 20 mg/mL DNase (Sigma) in

PBS for 1 hour at room temperature. The digest was filtered through 100 μ m nylon mesh to remove macroscopic debris and cell suspensions were washed and stained with antibodies. FACS sorting of tumor macrophages was performed as previously described (25, 46) using a BD FACSAria Cell Sorter to greater than 98% purity.

Preparation of bone marrow macrophages

Bone marrow cells isolated from long-bones of mice were cultured for a total of 7 days in complete media containing 40 ng/mL MCSF (Ebioscience), with additional growth media provided after 3 days of culture. Adherent cells were harvested and macrophage differentiation confirmed by flow cytometry for CD11b, F4/80, Gr1 and IA. Macrophages were differentiated into M1 or M2 phenotypes by culture for 24 hours in the presence of IFN γ (10 ng/mL Ebioscience) + 1 μ g/mL Ultrapure LPS (InvivoGen) or IL4 (10 ng/mL, Ebioscience), respectively as previously described (38), in the presence or absence of 1 μ g/mL or 25 μ g/mL RR-CDG.

Western blotting

Cells were lysed in RIPA buffer and denatured in SDS loading buffer containing β 2-mercaptoethanol, electrophoresed on 10% SDS-PAGE gels, and transferred to nitrocellulose. Blocked blots were probed overnight at 4°C with primary antibodies followed by HRP-conjugated secondary antibodies. Binding was detected using a Pierce SuperSignal Pico Chemiluminescent Substrate (Thermo Fisher Scientific) and exposure to film.

Statistical analysis

Data were analyzed and graphed using Prism (GraphPad Software). Individual datasets were compared using Student *t* test and analysis across multiple groups was performed using ANOVA with individual groups assessed using Tukey's comparison. A synergy evaluation for the endpoint of tumor size change between days 14 and 28 was performed using multivariable linear regression modeling, with RT treatment, CDG treatment, as well as their interaction term in the model. This analysis was performed using R 3.1.0 statistical program (R Foundation for Statistical Computing).

Results

Dose-dependent synergy between RR-CDG and RT

We have previously demonstrated that the tumor environment that emerges following RT suppresses adaptive immunity by activating a pathway of tissue repair and is driven by tumor macrophages (25, 38). We hypothesize that the potent inflammatory pathway activated by STING ligation will generate a strong adjuvant activity to enhance adaptive immune responses to antigens released by RT. To test the hypothesis, we evaluated synthetic CDN derivatives modified to be resistant to degradation by phosphodiesterases and to activate all known murine and human STING allele variants in a preclinical murine model of pancreatic cancer RT. We established Panc02 tumors in immune competent C57BL/6 mice and mice were randomized to receive radiation to the tumor, along with intratumoral injection of R_p, R_p dithio c-di-GMP (RR-CDG), or vehicle to that same tumor immediately following radiation and again at 24 hours (Fig. 1A, i). Radiation was delivered using CT-guidance via the SARRP and dosed to deliver a suboptimal dose of 10 Gy to the tumor with minimal

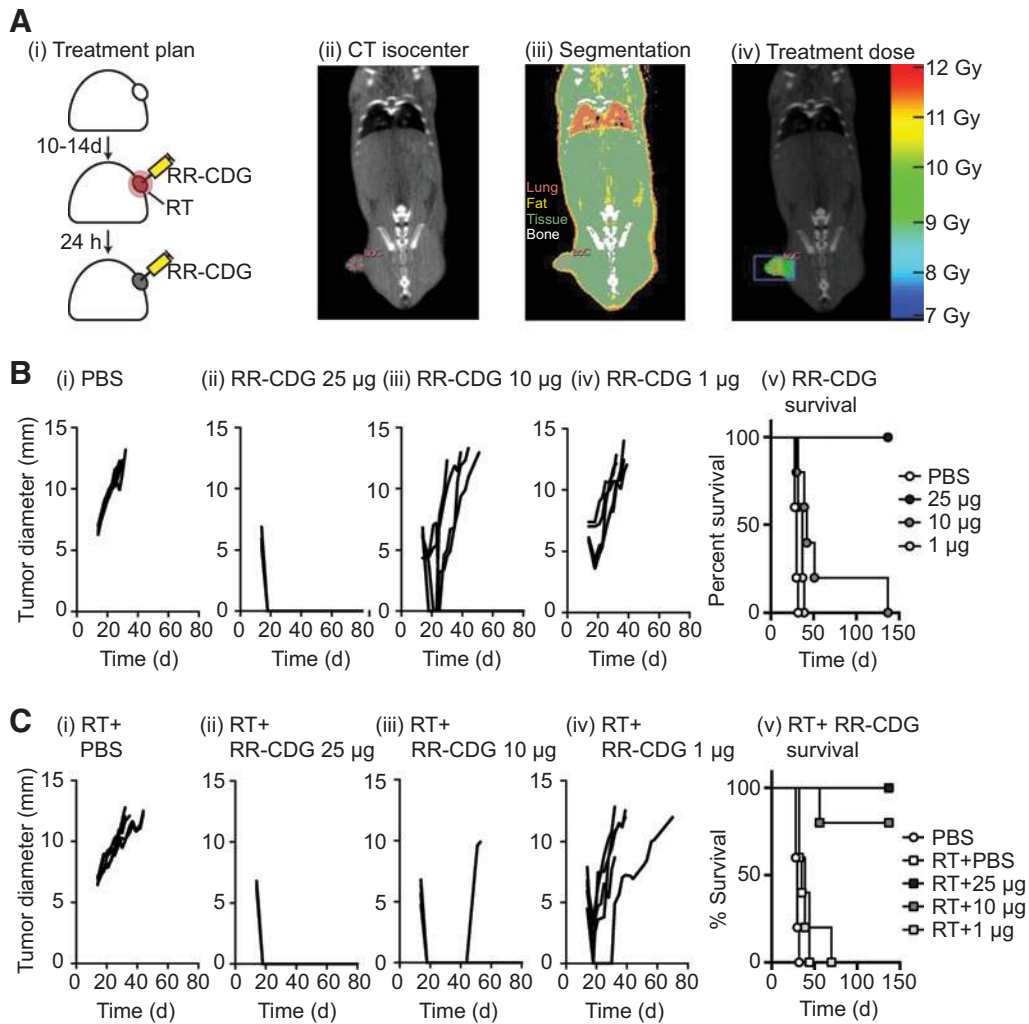


Figure 1. RR-CDG synergizes with RT for durable tumor cures. A, Panc02 pancreatic adenocarcinoma tumors were established in immune competent C57BL/6 mice and randomized to remain untreated or to receive RT immediately, followed by control (PBS) injections or 25, 10, or 1 μ g of RR-CDG into the tumor in matched volumes of PBS. ii, tumor isocenters were identified by CT guidance. iii and iv, CT images (iii) were used to determine tissue densities for dosimetry (iv) using the SARRP to deliver 10 Gy to isocenter with angled beams to minimize dose to radiosensitive organs. B, growth of individual tumors in mice not treated with RT (i-iv) and overall survival (v). C, growth of individual tumors in mice treated with RT (i-iv) and overall survival (v). Experiments incorporate 6 to 8 mice per group and the displayed experiment is representative of three independent repeats.

dose to surrounding normal tissues (Fig. 1A, ii-iv). RR-CDG was injected into the tumor at doses of 25, 10, or 1 μ g in PBS, or PBS vehicle alone and mice were followed for tumor growth and survival. RR-CDG caused a rapid, dose-dependent control of the tumor (day 4 following therapy PBS vs. 25 μ g RR-CDG $P < 0.001$; vs. 10 μ g RR-CDG $P < 0.001$; vs. 1 μ g RR-CDG $P < 0.01$), though at lower doses the tumor recurred (Fig. 1B). Of note, 10 Gy of RT alone did not significantly change tumor growth and minimally extended survival (median survival 30 days vs. 35 days, $P < 0.01$), but significantly increased survival with 10 μ g of RR-CDG (median survival 42 day vs. undefined, $P < 0.01$) and was lost at 1 μ g RR-CDG (median survival 37 days vs. 39 days; Fig. 1C). Similar responses were seen in mice bearing 3LL tumors treated with 25, 10, or 1 μ g RR-CDG or PBS along with 10-Gy RT (Supplementary Fig. S1). To determine whether these effects at the 10 μ g dose were synergistic, we tested for superadditivity using the endpoint of

tumor size change and found that the coefficient for RT:CDG was -4.11 , in the same direction with the coefficients for both RT and CDG, and a $P = 0.043$, indicating a super-additive effect of the combined therapy.

Adaptive immune control of distant tumors by RR-CDG plus RT

To determine whether this effect was a result of adaptive immune control of the tumor, the treatment was repeated in wild-type C57BL/6 mice as well as Rag1^{-/-} mice that lack adaptive immunity. Although combined therapy caused durable cure of tumors in wild-type mice, Rag1^{-/-} mice showed an initial tumor control but all recurred (Fig. 2A and Supplementary Fig. S2). To identify the cell type required, wild-type mice were depleted of CD8 T cells and similarly demonstrated initial control followed by recurrence. These data demonstrate that the durable cure of tumors by RR-CDG is composed of both an initial T-cell-

independent regression and long-term control dependent on CD8 T cells (Fig. 2A and Supplementary Fig. S2). To determine whether this therapy generated antigen-specific T cells, we repeated treatment in mice bearing Panc02 tumors expressing the model antigen SIY, and measured SIY peptide-specific T cells in the spleen 7 days following treatment. We demonstrate that only in mice receiving combination therapy were we able to detect significant increases in antigen specific cells in the spleen (Fig. 2B), suggesting this approach generates systemic T-cell immunity. To determine whether this would permit control of distant disease, we established a model where unmodified Panc02 tumors were established on opposite flanks, and only one tumor was treated with RT and RR-CDG (Fig. 2C, i). Only in mice where the tumor was treated with both therapies resulted in control of the distant tumor, though while there was a significant decrease in tumor size in these mice, not all mice in the treatment groups demonstrated benefit (Fig. 2C, ii). RT has been shown to attract antigen-specific T cells to the treatment site and can improve immune killing at the treatment site (3–5). To determine whether additional radiation of the uninjected tumor could enhance the effect, we repeated the experimental model with RR-CDG injection to only one tumor, but irradiated both tumors (Fig. 2D, i). In this setting, RR-CDG treatment resulted in a more dramatic control of the opposite flank tumor, and this did not occur in $Rag1^{-/-}$ mice or where CD8 T cells were depleted (Fig. 2D, ii),

demonstrating that the effect is immune mediated. These data demonstrate that radiation combined with RR-CDG generates systemic immune responses to tumor-associated antigens that can control distant disease. Panc02 tumors are spontaneously metastatic to the lung, and RT of the primary tumor alone does not affect metastatic tumor growth (Supplementary Fig. S3) as we have previously demonstrated with the 4T1 tumor model (45). Importantly, mice cleared of a single primary tumor by radiation plus RR-CDG did not develop metastatic disease over greater than 100 days of follow-up as determined by gross inspection and histologic analysis. These data demonstrate that radiation plus RR-CDG generates a functional systemic immunity to treated tumors *in vivo* that eliminates residual microscopic disease.

RR-CDG causes an early innate control by inducing stromal TNF α secretion

Because the initial tumor control by RR-CDG is not dependent on T cells, we investigated the mechanism of RR-CDG response in this early phase. Tumors harvested 6 hours following RR-CDG injection showed an extensive bloody infiltrate (Fig. 3A), and histologic evidence of early hemorrhagic necrosis (Fig. 3B). Agents such as the flavanoid DMXAA—also known as a vascular disrupting agent—which activates mouse but not human STING have been shown to have potent vascular effects (47), so we investigated whether this hemorrhagic necrosis was a result of

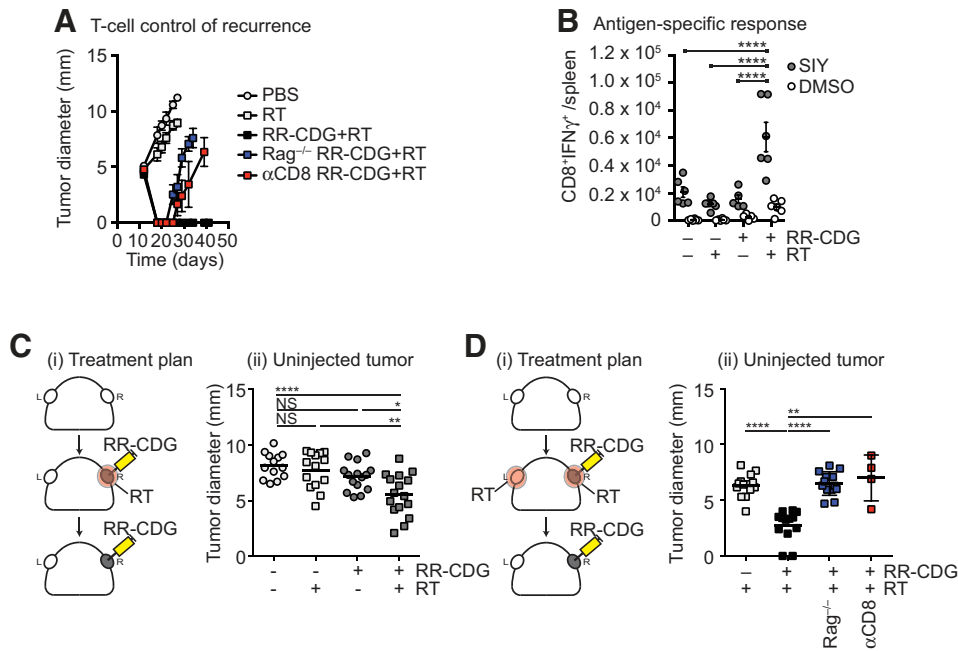


Figure 2. Radiation plus RR-CDG generates a T-cell response capable of controlling distant tumors. C57BL/6 or $Rag1^{-/-}$ mice challenged with Panc02 were left untreated or treated with 10 Gy focal radiation (RT) to the tumor, followed immediately by intratumoral injection of 25 μ g RR-CDG or PBS with a repeat injection the following day. Comparison groups of C57BL/6 mice were treated with anti-CD8 depleting antibodies 1 day before RT and again 1 week later. A, graphs show average tumor diameter. B, Panc02-SIY tumors were established and treated as in A; 7 days following treatment, spleens were harvested and tested for SIY-peptide-specific IFN γ production by intracellular cytokine staining. Graphs show the number of antigen-specific (SIY) or control (DMSO) CD8⁺IFN γ ⁺ cells per spleen. Each symbol represents one animal. C, i, duplicate Panc02 tumors were established on each flank of C57BL/6 mice and only one tumor was treated as in A. ii, graph shows tumor diameter of the untreated tumor 10 days following treatment. D, i, duplicate Panc02 tumors were established on each flank of C57BL/6 or $Rag1^{-/-}$ mice and both tumors were treated with 10-Gy RT, but only one tumor was treated with RR-CDG as in A. Additional groups were treated with anti-CD8 depleting antibodies 1 day before RT and again 1 week later. ii, graph shows the tumor diameter of the uninjected tumor 10 days following treatment. Each symbol represents one mouse. Statistics calculated by ANOVA with Tukey multiple comparisons test: *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.005$; ****, $P < 0.001$. The displayed experiment is representative of two independent repeats.

endothelial cell death. Immunohistochemistry for CD31 and smooth muscle actin demonstrated vascular cells remained in the tumor. However, these CD31⁺ cells lacked Smooth Muscle Actin⁺ (SMA⁺) podocyte coverage and were greatly enlarged (Fig. 3C). Immunostaining for F4/80 showed some clustering of macrophages around the enlarged immature vasculature (Fig. 3D), and there was no significant accumulation of CD3⁺ T cells in the tumor at this early stage. RR-CDG administration did not cause hemorrhagic necrosis when injected into tumor-free subcutaneous sites (not shown), and did not cause hemorrhagic necrosis when injected into the liver (not shown), which has extensive fenestrated vasculature, suggesting that the neoangiogenic vasculature of the tumor is particularly susceptible to RR-CDG treatment. To confirm these data in a more authentic model of pancreatic adenocarcinoma in mice, Pdx-Cre^{+/-} Kras^{(G12D)+/-}

Trp53^{(R172H)+/-} mice that spontaneously generate pancreatic tumors (42) were treated with RR-CDG administered to established pancreatic masses via laparotomy. Control tumors show invasive pancreatic adenocarcinoma impinging on residual normal regions of the pancreas (Fig. 3D, i). Twenty-four hours following RR-CDG administration, tumors show extensive hemorrhagic necrosis through the tumor stroma (Fig. 3D, ii). Interestingly, this hemorrhagic necrosis did not extend to adjacent normal regions of the pancreas (Fig. 3D, iii). The rapid induction of hemorrhagic necrosis was also demonstrated in the SCCVII head and neck cancer model on the C3H background, the spontaneous MMTV-PyMT mammary carcinoma model on the FVB background, and in the 3LL lung adenocarcinoma model on the C57BL/6 background (Supplementary Fig. S4). Though each model exhibited a different degree of hemorrhagic necrosis, each

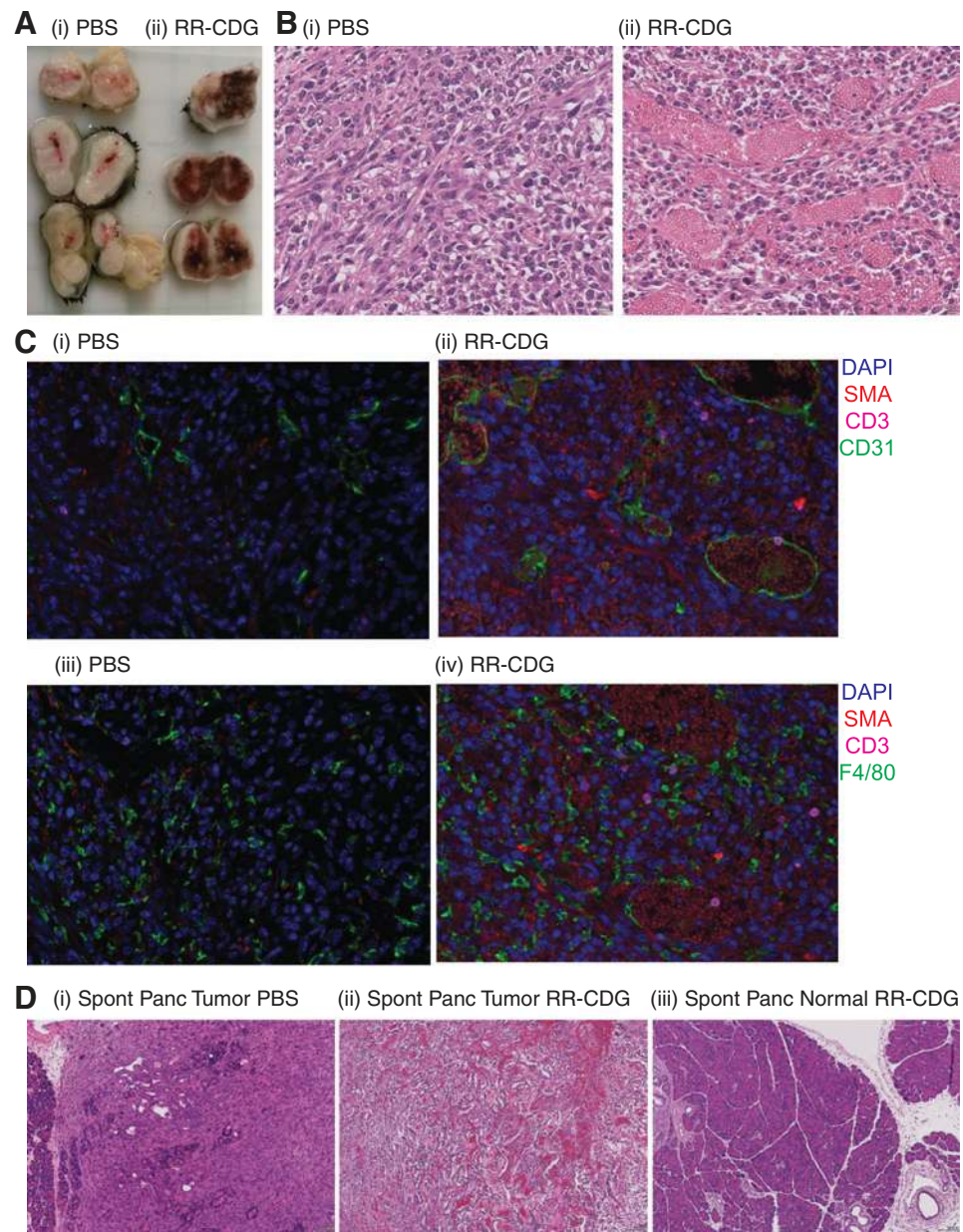


Figure 3. RR-CDG causes rapid vascular and macrophage reorganization in the tumor. A, established Panc02 tumors in C57BL/6 mice were treated with intratumoral injection of PBS vehicle (i) or 25 µg RR-CDG (ii) and harvested 6 hours later. At harvest, three RR-CDG-treated tumors showed macroscopic evidence of blood infiltrate. B, histology of control (i) and RR-CDG-treated (ii) tumors at 6 hours. C, immunofluorescence microscopy of control (i + iii) and RR-CDG-treated (ii + iv) tumors at 6 hours, with staining for smooth muscle actin (red); CD3 (magenta); a DAPI nuclear stain (blue) and either CD31 (green; i and ii) or F4/80 (green; iii and iv). Experiments are representative of two independent replicates that include four or more mice per group. D, Pdx-Cre^{+/-} Kras^{(G12D)+/-} Trp53^{(R172H)+/-} mice were treated with PBS (i) or RR-CDG (ii and iii) administered to established pancreatic masses via laparotomy. Images show representative sections of tumor (i and ii) or adjacent normal pancreas (iii).

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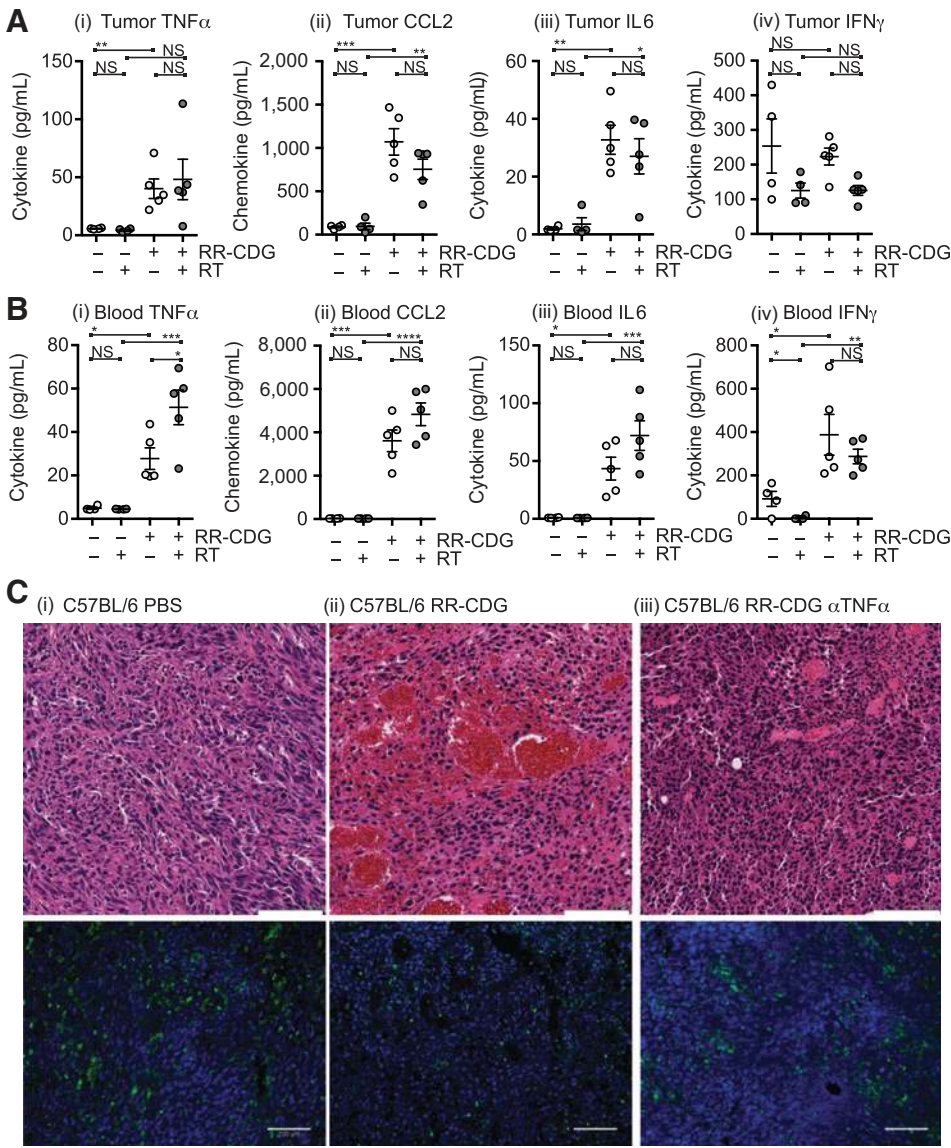


Figure 4. RR-CDG treatment results in cytokine secretion that causes early tumor regression. C57BL/6 mice challenged with Panc02 were left untreated or treated with 10 Gy focal radiation (RT) to the tumor, followed immediately by intratumoral injection of 25 μ g RR-CDG or PBS with a repeat injection the following day. Cytokine and chemokine levels were determined in the tumor (A) or peripheral blood (B) 6 hours following treatment by multiplex bead assay. Each symbol represents one mouse. Graphs additionally show mean plus SEM. C, C57BL/6 mice bearing established Panc02 tumors were treated with intratumoral injection of PBS (i), 25 μ g RR-CDG (ii), or 500 μ g anti-TNF α blocking antibody (iii) 6 hours before treatment with 25 μ g RR-CDG. Tumors were harvested 24 hours later for H&E staining in top images or immunostained for CD45 (green) alongside DAPI (blue) in bottom images. Scale bar, 100 μ mol/L. Statistics calculated by ANOVA with Tukey multiple comparisons test: *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.005$; ****, $P < 0.001$. Experiments incorporate 6 to 8 mice per group and the displayed experiment is representative of two independent repeats.

tumor showed a rapid decrease in size following treatment with RR-CDG and RT (Supplementary Fig. S4).

This early effect of RR-CDG resembles classic studies using TNF α injection into tumors (48). Analysis of cytokine and chemokine expression in tumors showed significant increases in TNF α , the monocyte chemoattractant CCL2, and IL6 at 6 hours following RT and RR-CDG treatment (Fig. 4A). This was accompanied by significant increases in these agents in the peripheral blood (Fig. 4B). Interestingly, IFN γ was significantly elevated in the peripheral blood by RR-CDG treatment but not in the tumor, suggesting that IFN γ secretion was stimulated outside the tumor environment. To determine whether RR-CDG mediated induction of TNF α was responsible for the early hemorrhagic necrosis in treated tumors mice were treated with anti-TNF α blocking antibody 6 hours before RR-CDG injection and tumors harvested 24 hours following RR-CDG injection. As expected, RR-CDG treatment caused hemorrhagic necrosis in the treated tumor, and this was largely blocked by injection of anti-TNF α blocking antibody

(Fig. 4C). Staining for CD45 $^{+}$ cells in these tumors demonstrated persistent immune cell infiltrate even in the presence of anti-TNF α blocking antibody (Fig. 4C).

RR-CDG acts through tumor stroma and repolarizes M2 tumor macrophages

To determine the cell type responsible for RR-CDG responses, tumors were stained for STING expression. Interestingly, both the cancer cells and the tumor stroma, including F4/80 $^{+}$ tumor macrophages, expressed STING (Fig. 5A). *In vitro*, both Panc02 cells and bone marrow macrophages respond to RR-CDG with upregulation of MHC1 and PDL1 (data not shown). To determine the relative importance of STING expression in the cancer cells versus the tumor stroma, Panc02 tumors were established in wild-type C57BL/6 mice or STING $^{-/-}$ (*goldenticket*) mice and treated with RT and RR-CDG. Interestingly, in STING $^{-/-}$ mice RR-CDG was entirely unable to cause early tumor regression, indicating that it is the stromal rather than cancer expression of STING that

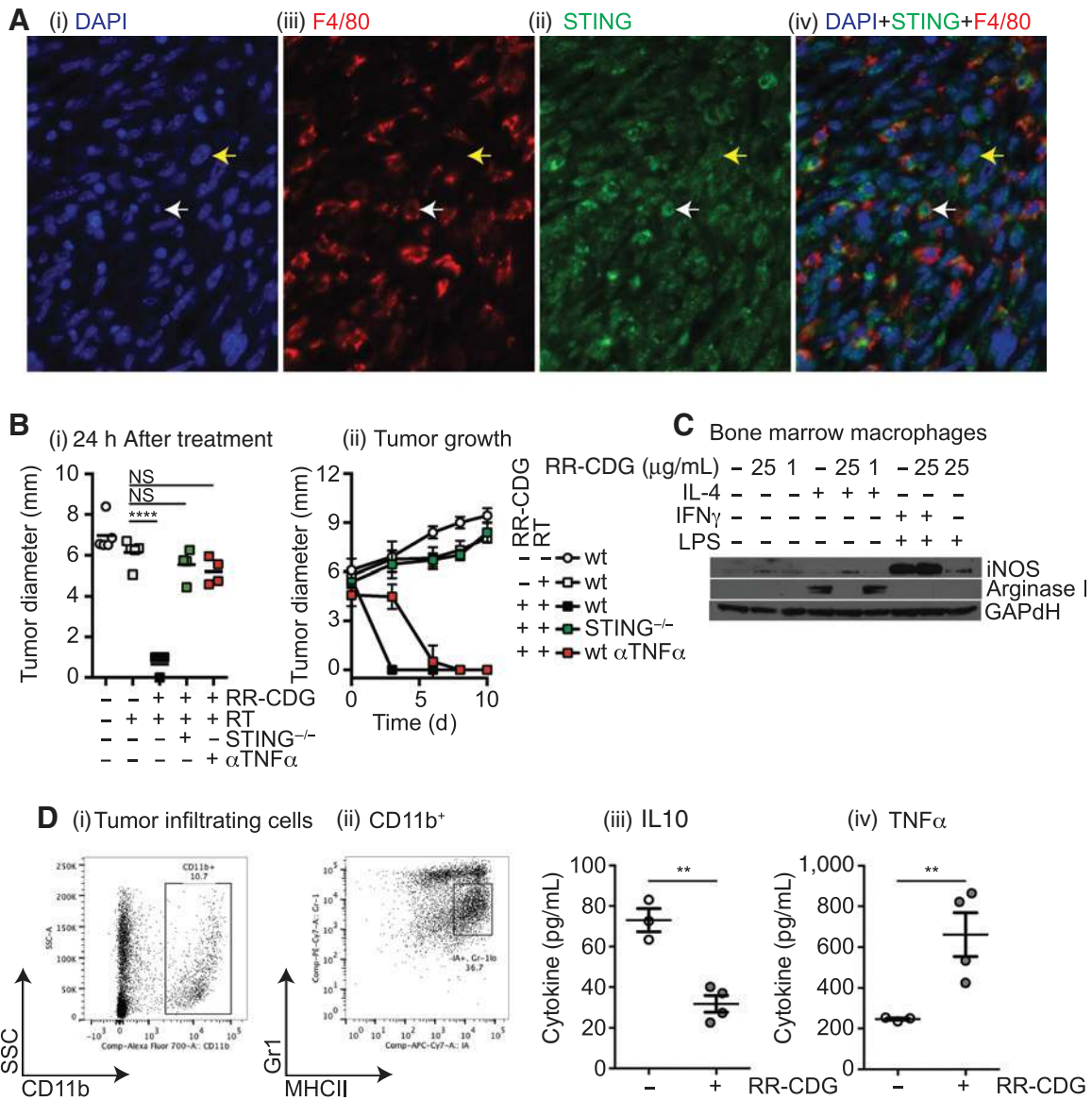


Figure 5. RR-CDG suppresses M2 differentiation and generates proinflammatory responses from tumor macrophages. A, immune histology in Panc02 tumors showing DAPI (i), F4/80 (ii), STING (iii), and a combined image (iv). White arrow, F4/80+ macrophage; yellow arrow, cancer cell. B, wild-type C57BL/6 or STING^{-/-} mice bearing established Panc02 tumors were left untreated or treated with 10 Gy focal radiation along with concurrent intratumoral injection of PBS or 25 μg RR-CDG and the injection was repeated 24 hours later. A control group of wild-type mice were treated with 500 μg anti-TNF α blocking antibody 6 hours before treatment with 25 μg RR-CDG and again 48 hours later. i, average tumor diameter 24 hours following treatment. ii, tumor growth over time following treatment. C, bone marrow macrophages were left untreated, treated with IL4 to direct M2 differentiation, or IFN γ and LPS to direct M1 differentiation in the presence or absence of RR-CDG at 1 or 25 $\mu\text{g/mL}$. Twenty-four hours later, macrophage differentiation was determined by Western blotting for iNOS and arginase I, with GAPdH as a loading control. D, sorted tumor macrophages were left untreated or treated with 25 $\mu\text{g/mL}$ RR-CDG for 6 hours, and secretion of IL10 (i) or TNF α (ii) was determined by bead assay. Statistics calculated by ANOVA with Tukey multiple comparisons test: *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.005$; ****, $P < 0.001$. Experiments incorporate 4 to 8 mice per group and the displayed experiment is representative of two independent repeats.

mediates this effect (Fig. 5B, i). Anti-TNF α also blocked the early regression caused by RR-CDG, in agreement with its ability to block hemorrhagic necrosis (Fig. 4C). Despite blocking early regression, tumors treated with anti-TNF α gradually regressed, while tumors in STING^{-/-} mice did not (Fig. 5B, ii), suggesting that even without early TNF α -mediated hemorrhagic necrosis,

RT combined with ligation of STING in tumor stroma results in tumor control through other immune-mediated control mechanisms.

We have previously demonstrated that macrophages accumulate in Panc02 tumors following RT and exhibit a suppressive M2 differentiation, characterized by expression of arginase I rather

than iNOS and secretion of IL10 and not TNF α following LPS stimulation (25). To determine whether RR-CDG could overcome this suppressive differentiation, we differentiated bone marrow-derived macrophages into a M2 phenotype by treatment with IL4, or into an M1 phenotype by treatment with IFN γ plus LPS. Concurrent treatment with 25 μ g/mL RR-CDG blocked arginase I upregulation by IL4, though was not able to cause significant iNOS expression when compared with the IFN γ plus LPS control (Fig. 5C). The effect was lost when the RR-CDG dose was dropped to 1 μ g/mL. These data suggest that RR-CDG treatment can block M2 differentiation of macrophages, though does not cause classic M1 differentiation. To determine the effect of RR-CDG on actual tumor macrophages, we purified tumor macrophages from Panc02 tumors (Fig. 5D, i and ii) and measured cytokine secretion following stimulation with RR-CDG. RR-CDG stimulation significantly decreased IL10 secretion by tumor macrophages and significantly increased TNF α (Fig. 5D, iii and iv) confirming that RR-CDG can overcome the suppressive differentiation of tumor macrophages, and that tumor macrophages are a potential candidate from the stromal response RR-CDG and a potential source of TNF β .

Pancreatic adenocarcinoma has STING expression in cancer cells and tumor stroma

The synthetic STING ligand RR-CDG is able to bind all isoforms of human STING (37). To determine whether pancreatic ductal adenocarcinoma has STING expression in a similar pattern to our murine model, we performed immunohistology for STING in patient samples following pancreaticoduodenectomy. We identified strong specific STING staining in cancer cells in examples of pancreatic ductal adenocarcinoma in patients that were treated with neoadjuvant chemoradiation before pancreaticoduodenectomy (Fig. 6A, i and ii). In these patients, residual pancreatic ductal adenocarcinoma cells as well as stromal cells express STING. Acinar cells in normal regions of the tumor-bearing pancreas do not express STING, though these normal regions still exhibit STING expression in stromal cells, and STING is detectable in both endothelial cells and ductal cells (Fig. 6A, iii) STING expression is also detectable in cancer cells in resection samples of untreated pancreatic adenocarcinoma (Fig. 6B, i). In contrast, normal duodenum epithelial cells neighboring the tumor resected during pancreaticoduodenectomy poorly express STING while underlying immune cells in the villi are strongly positive (Fig. 6B, ii). Similarly, the lymphoid aggregates occasionally visible in the stroma of pancreatic ductal adenocarcinoma show strong STING expression in a large proportion of cells (Fig. 6B, iii). These data demonstrate that human STING is present in stromal and cancer cells in pancreatic ductal adenocarcinoma and remains following chemoradiation, suggesting that these engineered cyclic dinucleotides targeting both murine and human STING in conjunction with RT are a potential novel combination for immunotherapy of pancreatic cancer.

Discussion

These data demonstrate that while radiation alone is poorly effective at generating systemic antigen specific immune responses, when combined with novel STING agonists, RT generates T-cell immunity required for local control and control of distant disease. Treatment with the synthetic RR-CDG ligand of STING resulted in a two-phase control of tumors.

First, there was a T-cell-independent hemorrhagic necrosis driven by TNF α secretion, followed by a CD8 T-cell-dependent control of recurrence. With higher doses of RR-CDG, the initial innate response is sufficient to control the tumor, but at lower doses where local control is incomplete, the ability of radiation to boost endogenous T-cell responses in the context of RR-CDG treatment results in T-cell-mediated control of both local and distant disease. RR-CDG treatment acts through cells in the tumor stroma, since its function is entirely dependent on STING expression by the host rather than the cancer cells, and we demonstrate that macrophages in the tumor are highly responsive to RR-CDG treatment.

RT has a great deal of potential to deliver site-directed cytotoxicity with minimal damage to nontarget tissues. The combined imaging technology, physics and computational dosimetry makes RT a rational intervention for therapy of individual lesions. Because radiation has rarely been able to affect tumors outside the treatment field, in patients with metastatic disease radiation has primarily been used as a palliative tool for individual problematic lesions. However, the recent combination of immunotherapy with radiation therapies has resulted in multiple reports of widespread metastatic tumor control in preclinical models and more recently in patients (15–18). Although the role of radiation in pancreatic cancer treatment has evoked controversy over the past two decades, the increasing study of stereotactic body RT in the neoadjuvant and locally advanced setting as reviewed by Franke and colleagues (49), provides an opportunity to use radiation in patients with borderline resectable and locally advanced pancreatic cancer both for its role in local control and as an immune modulator, alongside immunotherapy, to elicit systemic anti-tumor immune responses. As we gain a greater understanding of the immunological effects of RT, it is becoming clear that a complex interaction of multiple components is involved. First, radiation, at sufficient doses, kills cancer cells. The appropriate delivery and dose for cancer cell killing has been well established over decades of careful radiobiological research; however, the appropriate dose for priming immune effects is less clear. Fractionation of radiation over 5–7 weeks of daily treatment is a superior technique to ensure selective cancer versus normal cell death, though repeated treatment of the tumor likely causes repeated death of tumor infiltrating antigen specific T cells. Compacting radiation into fewer (1–5), higher doses of radiation made possible by advanced targeting techniques avoids continued treatments and is the approach used in almost all preclinical radiation models. However, we and others have shown that higher doses of radiation result in a wound-healing response involving tumor macrophages that likely suppresses adaptive immune responses at the treatment site (23–25). The second component brought by RT is a transient local improvement in the inflammatory environment of the tumor, resulting in cytokine and chemokine secretion and improved antigen presentation by cancer cells (3–5). Thus, when followed by adoptive transfer of strongly antigen-specific T cells, even low radiation doses can direct T cells to the tumor and improve local control where neither are effective alone (50). Together, these data suggest that where doses are high enough to kill cancer cells, RT can initiate new antigen specific immune responses if combined with immunotherapies sufficiently potent to overcome the poor context of antigen released by radiation. However, higher doses of RT are followed later by a wound-healing response that resolves adaptive immunity and limits immune control. The data reported here

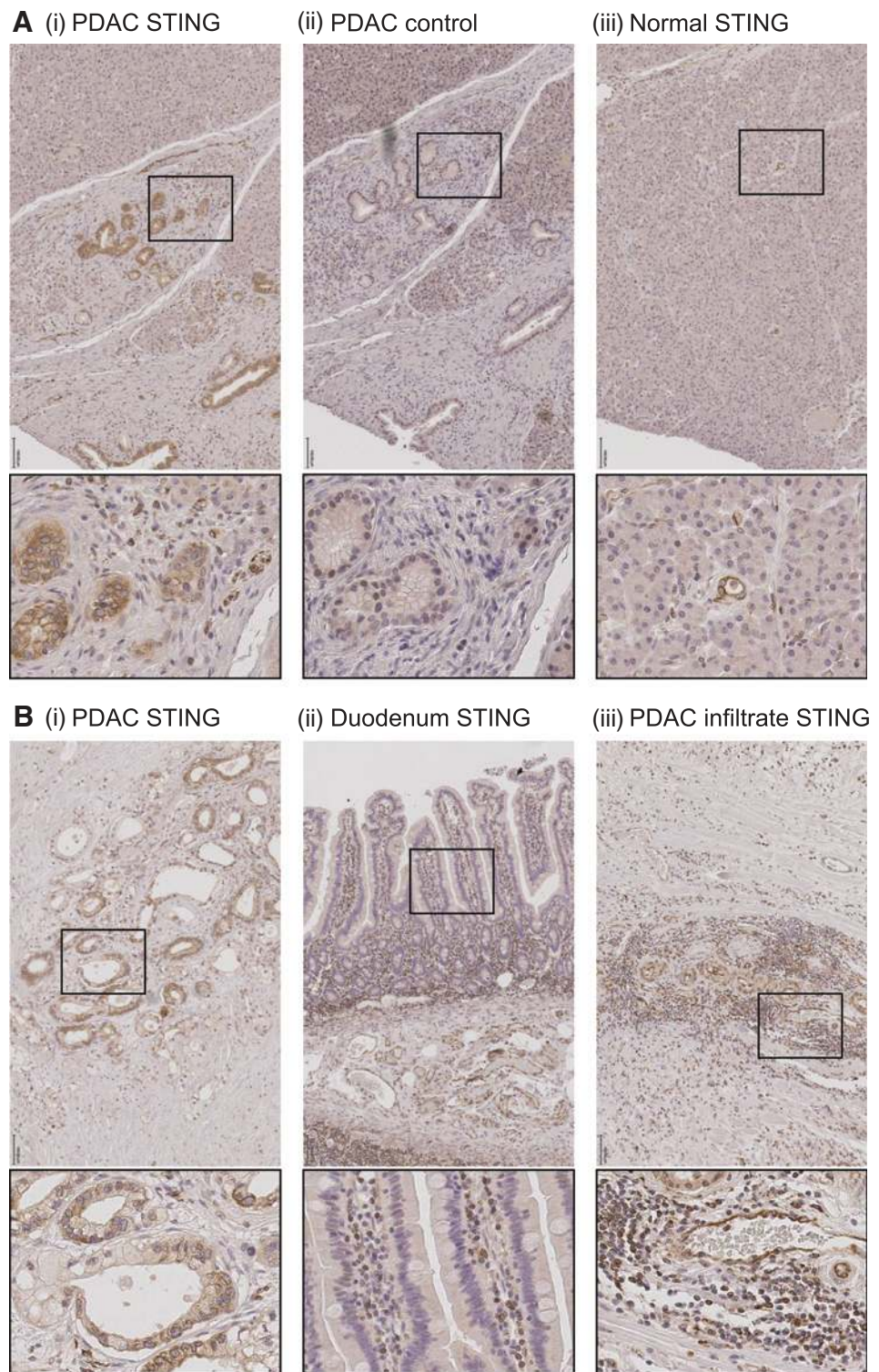


Figure 6. Expression of STING in pancreatic adenocarcinoma tumor and stroma. A, T3 pancreatic adenocarcinoma treated with neoadjuvant chemoradiation, followed by resection showing residual cancer (i and ii) or residual normal regions (iii) of the tumor-bearing pancreas stained for STING (i + iii) or control (ii) staining. B, resected tissue samples from untreated T3 pancreatic adenocarcinoma stained with STING (i); adjacent normal duodenum resected as part of pancreaticoduodenectomy (ii); and lymphoid aggregates in resected pancreas (iii). Scale bar, 100 μ mol/L. Insets show the origin of expanded images.

suggest that these novel ligands for STING can potentially improve the inflammatory context of antigens released by higher doses of RT and allow extended T cell control of residual disease.

Our data demonstrates that the rapid TNF α -driven hemorrhagic necrosis driven by RR-CDG is not necessary for tumor control, since early TNF α blockade limits this necrosis yet still results in

tumor regression. An alternative agonist of STING with reduced pyrogenic activity has been shown to have similar local control and improved ability to generate systemic T cell immunity as a single agent (37). Thus it is possible that the early TNF α -driven response limits T cell activity, potentially through an over-exuberant and possibly lymphotoxic cytokine response or a

suboptimal kinetic of antigen release and immune stimulation. This would fit with published reports where systemic administration of STING ligands can activate tolerogenic responses (51), as has been reported with CpG (52). This may explain why our data indicates that while T cells play a role in local tumor control with RR-CDG as a single agent there appears to be a failure to generate systemic immunity. It is possible that varying timing with regards to RT has the potential to improve the systemic adaptive immune response initiated by STING ligands. Nevertheless, the ability of STING ligation to generate a proinflammatory response even from macrophages polarized to suppressive phenotypes distinguishes this pathway from conventional adjuvants targeting Toll-like receptors (TLR). TLR ligation of macrophages purified from Panc02 tumors causes secretion of immune suppressive IL10 and not TNF α secretion in agreement with their M2 polarization (25), whereas we show here that STING ligation blocks IL10 secretion and generates TNF α . A range of agents have been shown to redirect polarization of macrophages, for example systemic administration of anti-CD40 generated more M1-polarized macrophages that were subsequently recruited to the tumor to improve pancreatic cancer therapy (53); however, few agents are able to generate M1-type responses from M2-type macrophages. The murine STING ligand DMXAA was recently shown to cause an early hemorrhagic necrosis in tumors that was dependent on macrophages and similarly repolarized macrophages from M2 to M1 phenotypes (47). In these experiments, it is interesting that tumors with higher macrophage infiltrates were more susceptible to DMXAA therapy (47) since commonly macrophages infiltrates causes resistance to T-cell-targeted immunotherapies such as anti-OX40, anti-CTLA4 and anti-PD1 (21, 54). These data suggest that therapies targeting STING may provide a novel treatment option for those patients most resistant to conventional immunotherapy.

Immunohistology shows that STING is well expressed in both the cancer cells and stroma of pancreatic tumors in patients, but is poorly expressed in normal regions of the pancreas. While expression of STING is required in stromal cells for tumor rejection by STING ligands, *in vitro* RR-CDG can cause upregulation of MHC I and PDL1 on STING-expressing cancer cells (data not shown) and non-hematopoietic cells can also self-sense DNA damage through STING (55). Thus, STING ligands have the potential to affect the cancer cells as well as the infiltrating cells as part of the desmo-

plastic reaction in pancreatic tumors. Nevertheless, since in our models RR-CDG function is entirely dependent on the expression of STING in stromal cells, the extensive stroma of pancreatic tumors remains an interesting independent target to accompany radiation-mediated death of cancer cells to generate a novel therapeutic combination.

Disclosure of Potential Conflicts of Interest

T.W. Dubensky, Jr. is the CSO of Aduro Biotech. M.R. Crittenden is a consultant/advisory board member for Regeneron. No potential conflicts of interest were disclosed by the other authors.

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Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): J.R. Baird, D. Friedman, B. Cottam, S. Bambina

Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): J.R. Baird, D. Friedman, B. Cottam, T.W. Dubensky, Jr., M.R. Crittenden, M.J. Gough

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Study supervision: M.J. Gough

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