

Overcome the challenge for intratumoral injection of STING agonist for pancreatic cancer by systemic administration Authors

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 cancer by systemic administration

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38

40 Abstract

Objective: Due to the challenge for intratumoral administration, innate agonists have not made it beyond preclinical studies for efficacy testing in most of tumor types. Pancreatic ductal adenocarcinoma (PDAC) has a T-cell excluded or deserted tumor microenvironment. Innate agonist treatments may serve as a T cell priming mechanism to sensitize PDACs to anti-PD-1 antibody (a-PD-1) treatment.

46 Design: Using a transplant murine model with spontaneously formed liver metastasis and 47 also the genetically engineered KPC mouse model that spontaneously develops PDAC, we 48 compared the antitumor efficacy between intrahepatic/intratumoral and intramuscular 49 systemic administration of BMS-986301, a next-generation STING agonist. Flow cytometry, 50 Nanostring, and cytokine assays were used to evaluate local and systemic immune responses.

51 Results: The study demonstrated that administration of STING agonist systemically via intramuscular injection is equivalent or potentially superior to its intratumoral injection in 52 inducing both effector T cell response and antitumor efficacy. Compared to intratumoral 53 54 administration, T cell exhaustion and immunosuppressive signals induced by systemic administration were attenuated. Nonetheless, either local or systemic treatment of STING 55 agonist was associated with increased expression of CTLA-4 in the tumors. However, the 56 combination of a-PD-1 and anti-CTLA-4 antibody with systemic STING agonist 57 demonstrated the antitumor efficacy in the KPC mouse spontaneous PDAC model. Our study 58 also demonstrated the feasibility and antitumor efficacy of systemic administration of BMS-59 986299, a new NLRP3 agonist. 60



62

Keywords: pancreatic ductal adenocarcinoma; stimulator of interferon genes; innate immune
agonist; STING; NLRP3; tumor microenvironment; immune checkpoint inhibitor;
immunotherapy; systemic administration; intratumoral injection.

via systemic administration, instead of local administration, for treating PDAC.

67

What is already known on this topic – Despite promising preclinical studies, innate
immune agonists including STING agonists and NLRP3 agonists have not been tested in
most of tumor types due to the difficulties associated with their intratumoral delivery.

71 What this study adds – This study compared between intratumoral and systemic 72 administration of BMS-986301, a next-generation STING agonist and BMS-986299, a 73 new NLRP3 agonist in the preclinical models of pancreatic cancer. Notably, systemic 74 administration of STING agonist showed comparable or potentially superior effector T cell 75 response and antitumor efficacy compared to intratumoral administration, with attenuated T 76 cell exhaustion and immunosuppressive signals.

How this study might affect research, practice or policy – For the first time, this study
supports the clinical development of innate agonists via systemic administration for treating
pancreatic cancer. Supported by the results in this study, a phase 1 trial evaluating BMS986301 intratumoral or systemic intravenous injection as monotherapy or in combination
with nivolumab (PD-1 blockade) and ipilimumab (CTLA-4 blockade) in patients with
advanced solid cancers has been initiated (NCT03956680).

84 Introduction

Recent research has highlighted the crucial role of the innate immune system in tumor 85 immunosurveillance and the stimulation of antitumor immune responses[1]. This has led to 86 87 the development of several small-molecule innate agonists as potential immunotherapeutics 88 or vaccine adjuvants for different types of cancer[2]. However, despite promising preclinical studies, these agents have not been tested in most of tumor types due to the difficulties 89 associated with their intratumoral delivery[3]. Overcoming this challenge is critical for the 90 clinical development of innate agonists. Nonetheless, the potential benefits of using innate 91 92 agonists as T cell priming agents continue to provide an impetus for the exploration of novel strategies to overcome the challenge for their delivery. 93

94

95 Stimulator of interferon genes(STING) is a transmembrane protein that is expressed in various endothelial, epithelial and hematopoietic cells. Upon activation at the endoplasmic 96 reticulum and subsequent translocation to the Golgi, STING recruits and activates TANK-97 binding kinase 1(TBK1), which in turn phosphorylates and activates interferon regulatory 98 factor 3(IRF3) and NF-kB transcriptional programs, resulting in the expression and release of 99 100 pro-inflammatory type I interferons(IFNs) and cytokines[4, 5, 6]. Accumulating evidence has suggested that STING also possesses cell-intrinsic tumor suppressive activity[7]. The 101 mechanistic underpinnings of the cyclic GMP-AMP synthase-STING pathway make STING 102 agonists(STING-A) a promising adjuvant to cancer vaccines[8]. Numerous natural or 103 104 synthetic STING-A as monotherapy or in combined with other treatments have also been tested in both pre-clinical studies and clinical trials across many cancer types[9, 10]. 105

Synthetic cyclic dinucleotides(CDNs) were the first generation of STING-A that entered the
clinical trial phase of drug development due to their structural versatility and ability to bind
all prevalent allelic variants of human STING.

109

110 Pancreatic ductal adenocarcinoma(PDAC) is one of the most lethal malignancies due to its 111 resistance to conventional therapies. PDAC and other non-immunogenic "cold" tumors do not respond to immune checkpoint inhibitors(ICIs) as monotherapy largely due to the lack of 112 113 tumor-infiltrating effector lymphocytes [11, 12]. Published studies have demonstrated that 114 intratumorally injected STING-A inflamed the tumor microenvironment(TME) of PDAC with an effector T cell infiltration and reduced tumor burden in the mouse model of 115 PDAC[13]. However, despite these promising results at the preclinical phase, anti-tumor 116 117 efficacies of STING-A have not been substantiated, largely due to the challenge of intratumoral delivery of such agents, which may have never reached therapeutic dose levels 118 or never been tested in the appropriate disease indications. 119

120

BMS-986301 is a novel CDN-based, next-generation STING-A and has demonstrated a promising antitumor activity in the CT26 and MC38 subcutaneous tumor murine models, resulting in more than 90% of tumor regression compared to only 13% with the firstgeneration STING-A, ADU-S100. Similar results were observed with a single intratumoral administration of BMS-986301 in combination with an anti-PD-1 agent [14]. However, these previous studies were limited by the subcutaneously implanted tumors, which do not resemble the TME of human PDAC[15]. Therefore, this present study aims to deliver this next-generation CDN-based STING-A systemically and compare its anti-tumor efficacy and
elicited immune response with the intratumoral injection of this agent in a liver metastasis
model of PDAC.

131

132 Methods

133 Mouse experiments

The mouse study was conducted in accordance with the guidelines established by the Animal Care and Use Committee of Johns Hopkins University. Female C57Bl/6 mice aged 6 to 8 weeks were purchased from Jackson Laboratories and maintained under the Institutional Animal Care and Use Committee(IACUC) guidelines. Third-party management was responsible for maintaining the IACUC mouse protocol.

139

STING-A(BMS986301, BMS) was dissolved into DPBS(Life Technologies) vehicle and
administered to tumor-bearing mice either by intratumoral or intramuscular injection once a
week at a dose of 5mg/kg starting on day 14, for a total of three doses. a-PD-1(BMS936558,
BMS) and an IgG control(ab18443, BMS) were administered intraperitoneally twice weekly
starting on day 14, for a total of five doses, at a dose of 10mg/kg.

145

146 NanoString

147 After the mouse was euthanized and the tumor was harvested, the tumor tissues were 148 submerged into RNA-later(Invitrogen) to preserve the RNA. The total RNA was extracted 149 from the whole specimen using the AllPrep DNA/RNA/Protein Mini Kit(Qiagen) according

100	to the manufacturer's instructions. The RNA was then equalized for NanoString hybridization
151	using the Formulatrix Tempest. The murine PanCancer Immune panel codeset
152	XT_PGX_MmV1_CancerImm_CSO, Cat#115000142), which contains 750 target genes
153	along with housekeeping and negative/positive control probes, was used for the NanoString
154	hybridization. The data obtained was then analyzed using NanoString nSolver 3.0 and an
155	internally developed NanoString Data Analyzer Rshiny app(BMS). The samples were run on
156	the NanoString MAX system reader, and the data was analyzed using various CRAN and
157	Bioconductor packages such as dplyr, tidyr, and reshape2 to clean, reformat and match the
158	sample annotations to the normalized data exported from NanoString nSolverV.3.0.
159	
160	Additional methods are provided in the Supplemental Materials.
161	
162	Results
162 163	Results Develop a mouse model to resemble the intratumoral injection of STING-A in
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die from liver metastases if not treated[17]. The changes in the sizes of the metastatic lesions
in the liver can be monitored by small animal ultrasonography. Liver metastases have been
identified as potential sites for the IT administration of the first-generation STING-A to treat
patients with PDAC. In this study, the liver metastases serve as the clinically relevant target
tumors for IT administration of STING-A(Fig.1A).

177

Mice were also inoculated subcutaneously with the KPC tumor cell line to form 178 subcutaneous(SubQ) tumors, not for the IT injection of STING-A, but for the abscopal effect 179 180 to be examined. As shown in Fig.1B, sixteen mice per group were treated by control vehicles(negative control, NC), a-PD-1, STING-A, and STING-A+a-PD-1(Combo). A total 181 of five doses of a-PD-1 was administered twice a week by intraperitoneal injection(IP), while 182 183 the STING-A was given for three weekly doses by IT injection to mimic local treatment or intramuscular(IM) injection to mimic systemic treatment. Mouse survival was followed; the 184 target liver metastatic lesion was measured by ultrasound; and the subQ tumors was 185 186 measured by calipers twice a week. We did not observe any obvious sign of toxicity including bleeding, infection, paralysis, and weight loss as a result of IT injection of STING-A. 187 188 However, we noticed small areas of liver necrosis around the injection sites at the necropsy of two mice(Fig.S1B). It should be noted that only approximately 35% of mice in the 189 hemisplenectomy model harbored a liver metastasis feasible for IT injection. 190

191

192 The results(Fig.1C) showed that the tumor growth inhibition(TGI) rate of the target liver 193 metastasis lesion was significantly increased in the STING-A and a-PD-1 combination

194 treatment group(maximum TGI=88.68±98.42%, p <0.05) and the STING-A group(maximum TGI=69.38±73.94%, p <0.05) as compared to the negative vehicle group. To investigate 195 196 whether the local intratumoral injection of STING-A could induce the abscopal effect, we implanted subcutaneous tumors on the bilateral flanks of the liver metastasis mice model to 197 mimic distant metastases. The TGI rate of bilateral SubQ tumors was also significantly 198 increased in the combo group(maximum TGI=66.38±86.00%, p <0.05) as compared with 199 200 negative control(Fig.1D). Interestingly, the TGI in the distant SubQ tumors is bigger than that of the locally targeted liver metastatic lesion, supporting an abscopal effect from the STING-201 202 A and a-PD-1 combo treatment. Moreover, we compared the survival of liver metastasis mice 203 in the above treatment groups and found that the combo treatment significantly prolonged survival comparing to the control treatment and the STING-A treatment(Fig.1E). 204 Nevertheless, the combo treatment prolonged survival modestly without a statistical 205 significance comparing to the a-PD-1 treatment. It is possible that the local intratumoral 206 injection of saline may have caused inflammatory response which resulted in a small effect 207 on priming the tumor for the a-PD-1 treatment. Nevertheless, the combo treatment 208 significantly prolonged survival comparing to the control treatment and the STING-A 209 210 treatment(Fig.1E).

211

STING-A in combination with anti-PD-1 antibody enhances effector T cells infiltration
and CD103⁺dendritic cells in both target liver metastatic lesions and non-target
metastatic lesions

215 To understand the mechanistic basis of the enhanced anti-tumor activity of STING-A in

216 combination with a-PD-1, we performed flow cytometry analysis of the tumor infiltrating leucocytes(TILs) derived from the targeted liver metastasis in the same PDAC mouse model. 217 218 As described in the dosing schema in Fig.1B, mice received the a-PD-1 on days 14 and 17, and the STING-A on days 14. On day 21, the mice were sacrificed and the implanted SubQ 219 220 tumors as well as livers were harvested. TILs from the targeted liver metastatic lesion and the 221 whole liver tissue excluding the targeted lesion were compared. Note that the whole liver 222 tissue harvested on day 21 was diffusely infiltrated by non-target metastatic lesions. The results showed that the STING-A+a-PD-1 combo therapy significantly increased the 223 infiltration of the CD8⁺ and CD8⁺PD-1⁺T cells, but not the CD4⁺ and CD4⁺PD-1⁺T cells in 224 225 the target lesion comparing to treatment controls(Fig.2A;Fig.S1C). Interestingly, a-PD-1, STING-A, or their combo all decreased the MHCII⁺CD11c⁺dendritic cells(DCs); however, 226 227 comparing to the treatment control, a-PD-1 alone, or STING-A alone, the a-PD-1 and STING-A combo significantly increased the CD11b⁻CD103⁺ subtype of DCs(Fig.2B), which 228 are known to play a role in the cross-presentation of tumor antigens[20]. TILs from non-229 target liver metastases showed that the combo treatment resulted in a significant increase in 230 CD8⁺T cells comparing not only to the treatment control, but also to a-PD-1 alone(Fig.2C). 231 232 The infiltration of CD8⁺PD-1⁺T cells in the combo group was comparable with the control group and significantly lower than that in the STING-A alone group(Fig.S1D). Moreover, 233 CD4⁺T cells and CD4⁺PD-1⁺T cells were both significantly decreased in the combo treatment 234 group compared to the STING-A treatment group, presumably due to the treatment effect of 235 a-PD-1(Fig.2C). These results suggest that, in non-target liver metastatic lesions, CD8⁺ and 236 CD4⁺T cells both trended in the favor of anti-tumor immune response following the a-PD-1 237

238 and STING-A combo treatment. In non-target liver metastases, MHCII⁺CD11c⁺DCs were similar among all treatment groups, suggesting that an enhanced antigen presentation was 239 originated in locally targeted lesions. However, CD11b⁻CD103⁺DCs were significantly 240 elevated in the combo treatment group in non-target lesions(Fig.2D). It is possible that 241 242 CD103⁺DCs trafficked from targeted lesions to non-target lesions. Taken together, these 243 results suggest that both STING-A and a-PD-1 are required to activate local immune response in favor of anti-tumor response and that this immune response is extended to the non-target 244 lesions in the vicinity of the target lesion. 245

246

STING-A in combination with anti-PD-1 antibody activate anti-tumor immunity via innate immune response signaling pathways

249 As the flow cytometry analysis has limitation in examining cytokine/chemokines and 250 intracellular signals involved in the innate immune response, we examined the gene regulation of innate immune response following STING-A and/or a-PD-1 treatment by using 251 the NanoString assays with the murine PanCancer Immune panel. We selected differentially 252 expressed genes by comparing the STING-A and a-PD-1 combo treatment and treatment 253 254 controls, STING-A alone or a-PD-1 alone with a 5% false discovery rate(Fig.S1E). We first examined the gene expression in the interferon(IFN) response pathways. As shown in Fig.2E, 255 256 the gene expression of IFN- γ (*Ifng*) was significantly increased in the combo treatment group and was also accompanied by significantly increased expression of Ifngr1, Ifnar2, Irf1, Irf4, 257 Irf5, Irf7, Irf8, Stat1, Stat2, Stat4, Ifitm1, Ifih1, Ifit1, Ifit2, Ifit3, Ifi35, and Ifi44 compared to 258 the control or single treatment groups. However, Irf3 was significantly decreased in the 259

260 combo treatment group compared to the a-PD-1 treatment group(Fig.S1F). In addition, the tumor necrosis factor(TNF)-response pathway genes including *Tnf*(encoding TNF-α), 261 Lta(encoding TNF-B), Nfkb1, Nfkb2, and Tank were also significantly upregulated in the 262 combo treatment group compared to the control or single treatment groups(Fig.2F). The 263 264 innate immune response pathways genes including Tmem173(STING), Ddx58, Nlrp3, Nlrc5, 265 Myd88, Clec4a2, Clec4n, Clec5a, Clec7a, Nod1, and Nod2 were significantly increased in the 266 combo treatment group compared to the control or single treatment groups(Fig.2G). However, as anticipated, the gene expression of Jak1 and Jak2 was similar among all treatment groups, 267 268 suggesting Jak1/2 are not regulated at the RNA level(Fig.S1F).

269

We next investigated the expression of cytokines which are known to be involved in the 270 271 development and differentiation of T and B lymphocytes(Fig.3A-B;Fig.S2A-B). As shown in 272 Fig.3A, the expression of most genes in the interleukin(IL)-1 and IL-18 family except *Il18* 273 itself and *Irak1* was significantly increased in the combo treatment group compared to most of other treatment groups. As demonstrated in Fig.3B, the gene expression of most of the 274 proinflammatory cytokine receptors including Il2ra, Il2rb, Il2rg, Il7r, Il12rb1, Il12rb2, 275 276 113ra1, 1113ra2, and 1115ra was significantly increased in the combo treatment group 277 compared to most of other treatment groups. Interestingly, the cytokines themselves including Il2, Il7, Il12, and Il13, besides Il21 which was significantly upregulated, were upregulated in 278 279 a non-statistically significant trend in the combo treatment group compared to the control or single treatment groups(Fig.S2A). These results suggest that pro-inflammatory pathways 280 including those that mediate the inflammasome are activated broadly by the combination of 281

As anticipated, we found that the majority of chemokine genes were upregulated in the 284 combo treatment group compared to the a-PD-1 treatment group(Fig.S2C), likely due to the 285 286 innate immune response induced by the STING-A. Therefore, we focused on those associated 287 with T cell trafficking. Our results demonstrate a significantly increased expression of *Ccl1*, 288 Ccl2, Ccl3, Ccl4, Ccl5, Ccl7, and Ccl8 in the combo treatment group compared to the vehicle control treatment group(Fig.3C), suggesting that stimulation of innate immune response is 289 290 anticipated to induce myeloid cell infiltration. C-X-C motif chemokine ligand(CXCL) 9, CXCL10, and CXCL11 are known to bind C-X-C motif chemokine receptor(CXCR) 3 on T 291 cells and, in response to IFN signaling to recruit memory and activated effector T cells[21]. 292 293 We observed a significant enhancement of the expression of Cxcl9, Cxcl10, and Cxcl11 in the combo treatment group compared with the control or single treatment groups(Fig.3C) 294 although the expression of Cxcr3 was similar among different treatment groups(Fig.S1E). 295 296 Interestingly, comparing to the vehicle control treatment group, the administration of a-PD-1 showed a statistically non-significant trend of increase whereas STING-A showed a trend of 297 298 decease in the expression of *Ccl17*, which encodes a T regulatory cell(Treg) chemokine. This finding is thus consistent with published studies showing that ICIs upregulate C-C motif 299 chemokine ligand(CCL) 17 expression in tumors and increase the migration of Tregs into the 300 TME of PDAC[22, 23]. Moreover, we here observed that the combo treatment led to a 301 significantly decreased expression of Ccl17(Fig.3C;Fig.S2D). It should be noted that the gene 302 expression results from the NanoString assay may be influenced by an influx of immune cells 303

that express the genes. Therefore, an increased expression of certain immune genes may represent an increased infiltration of the relevant immune subtypes. Taken together, these results suggest that STING-A may confer an antitumor effect by suppressing CCL17 expression or CCL17-expressing cells and thereby suppressing Treg migration into the TME.

308

309 STING-A in combination with anti-PD-1 antibody enhances T cell activation signals

310 Among those differentially expressed genes in the above NanoString analysis, we did further statistical analysis on genes related to the activation of effector T cells. As shown in Fig.S2E 311 312 and as anticipated, the expression of T cell receptor/CD3 signaling pathway genes including Cd3d, Cd3e, Cd3g, Cd3z, Cd8a, and Lck in the combo treatment group was significantly 313 increased[24, 25]. In addition, our results(Fig.3D) revealed that Gzma, Gzmb, Gzmk, and Prf1 314 315 expression was significantly elevated in the combo treatment group comparing to most of other treatment groups, suggesting that the cytotoxic function of effector T cells are 316 significantly enhanced. The gene expression of T cell co-stimulatory factors including 317 Tnfrsf9(CD137), Tnfrsf4(OX40), and Icos were significantly increased in the combo 318 treatment group when compared to any other group and including Cd27 when compared to 319 320 the vehicle treatment group(Fig.3E). However, the gene expression of co-inhibitory receptors including Pdcd1(PD-1), Lag3, Havcr2(TIM3), and Ctla4 and the expression of immune 321 checkpoint activators such as Cd274(PD-L1) and Ido1 were significantly elevated in the 322 combo treatment group compared to any other treatment group(Fig.3F;Fig.S2F). These 323 324 results suggested that T cell activation in the combo treatment group may also lead to the T cell exhaustion, in consistence with previously published studies[26]. 325

327 STING-A in combination with anti-PD-1 antibody enhances the infiltration and 328 activation of effector T cells in the distant subcutaneous tumors

329 We next examined the immune response that mediates the abscopal effects in the distant 330 tumors. As shown in Fig.4A, we observed a significant increase in the infiltration of $CD8^+$ 331 and CD8⁺PD-1⁺T cells in the SubQ tumors from the combo treatment group, supporting the abscopal effect. Similar to the locally targeted lesions, CD4⁺ and CD4⁺PD-1⁺T cells in the 332 333 SubO tumors were not significantly changed among all treatment groups. 334 CD11b⁻CD103⁺DCs in the SubQ tumors were also similar among all treatment groups, suggesting that the activation of antigen presenting cells occurred locally(Fig.4B). 335

336

337 We then used the same NanoString assay to assess the T cell functional status. The results indicate that the expression of genes associated with the T-cell receptor CD3 complex 338 exhibited a similarly significant increase in the SubQ tumors from the combo treatment group 339 compared to other treatment groups as in the liver metastases(Fig.S3A). Similarly, genes 340 related to the cytotoxic activities of effector T cells, including Gzma, Gzmb, Gzmk, and Prf1, 341 342 demonstrated a significant increase in the combo treatment group compared to most of other treatment groups(Fig.4C). In addition, the combo treatment group exhibited a significant 343 increase in the expression of signals related to T cell activation including Cd27, Icos, 344 Cd274(PD-L1), and Idol comparing to most of other treatment groups(Fig.4D-E) and 345 including Cd28, Cd80, Pdcd1(PD-1), Lag3, and Ctla4 comparing to the vehicle treatment 346 group(Fig.S3B). Consistently, genes related to chemokines for effector T cell trafficking 347

348 including Cxcl9, Cxcl10, and Cxcl11 exhibited a significant increase in their expression in the combo treatment group compared to most of other treatment groups(Fig.4F). Ccl17 also 349 350 exhibited a significant decrease in the combo treatment group compared to the vehicle treatment group. Interestingly, the expression of *Ccl28*, a T and B cell homing factor[27], 351 352 showed a significant increase in the combo treatment group in SubQ tumors(Fig.4F), but not 353 the target metastatic lesions(Fig.S2C). The gene expression of cytokines and cytokine receptors that are relevant to the activated status of T cells was similarly increased as seen in 354 355 the liver metastases, again except *Il18*(Fig.4F), further supporting the abscopal effect.

356

In addition, the profiles of other cytokines and chemokines in the SubQ tumors were similar 357 to those in the targeted liver metastases and showed a significant increase in pro-358 359 inflammatory immune responses in the IFN(Fig.4G) and TNF pathways(Fig.4H), but no significant changes or a decrease in Jak1 and Jak2(Fig.S3C), in the combo treatment group 360 compared to other treatment groups. Nevertheless, expression of innate agonist receptors and 361 adaptors in SubQ tumors appeared to be somewhat different from that in targeted liver 362 metastases, showing an increase in *Tlr8*, *Nlrc5*, *Nlrp3*, *Clec4n*, *Clec5a*, *Clec7a*, and *Nod1*, but 363 364 a decrease in *Ticam1* and *Mavs* in the combo treatment group compared to other treatment groups(Fig.4I). However, the profile of chemokines and chemokine receptors that function in 365 the myeloid cell trafficking was similar between targeted liver metastases and distant SubQ 366 tumors in the combo treatment group compared to other treatment groups(Fig.S3D). 367

368

369 Establish a mouse model for systemic administration of STING-A in combination with

370 anti-PD-1 antibody induced both systemic and intratumoral immune responses

We next tested the IM administration, a systemic administrative route, of STING-A. To 371 372 compare the anti-tumor efficacy of IM with IT injection of STING-A, we used the same liver metastasis model with implantation of SubO tumors by following the same schema described 373 374 in Fig.1B. Firstly, systemic administration did not show any obvious toxicity including 375 bleeding, unhealed wound, paralysis, or weight loss, etc. All the mice following the 376 hemisplenectomy procedure were candidates for IM injection although we chose those feasible for IT injection for the purpose of comparison. We then found that mice in the 377 378 combo(IM) group had a significantly prolonged survival when compared to other treatment groups(Fig.5A). The TGI rate for SubQ tumors also exhibited a significant increase in the 379 combo(IM) group(maximum TGI=58.86 \pm 49.12%, p <0.05)(Fig.5B). By combining the 380 381 survival data in both IT and IM experiments, we found that the majority of mice died at around 3 to 4 weeks after tumor implantation whereas those who received both STING-A by 382 IT or IM and a-PD-1 may live up to 6 weeks(Fig.S4A). Mice who received STING-A 383 intramuscularly in combination with a-PD-1 reached the longest survival beyond 6 weeks 384 although, likely due to the small sample size, there was no significant survival difference 385 386 between the IM combo and the IT combo group.

387

To assess the systemic immune response induced by the IM injection of STING-A and/or a-PD-1, we first measured the cytokine response in the serum samples harvested 6 hours after the IM injection of STING-A. The results demonstrated that a number of cytokines especially those associated with inflammation had a significantly increased level in the sera of mice 392 from the combo treatment group, including TNF- α , and IFN- γ (Fig.5C). Interestingly, several T lymphocytes trafficking chemokines including CXCL9, and CXCL10 and type I cytokines 393 including IL-2[28] and IL-12[29] were significantly increased in the sera from mice in the 394 395 combo treatment group compared to other treatment groups, suggesting that IM injection of STING-A in combination with a-PD-1 is potentially able to induce an anti-tumor systemic 396 397 immune response(Fig.5D). In addition, some interleukins including IL-5[30], IL-6[31], and 398 IL-10[32] that support lymphocyte growth and/or antibody production were also boosted in the IM combo group(Fig.S4B). The results also indicated the increased production of 399 400 cytokines that participate in the recruitment of macrophages, neutrophils, and eosinophils, including CCL2, CCL3, CCL4, CCL5, CXCL1, granulocyte colony stimulating factor(G-401 CSF), and CCL11(Fig.S4C). These results suggested that the IM injection of STING-A is 402 403 able to induce similar systemic immune responses as evidenced by the above described Nanostring analysis following the IT injection of STING-A. 404

405

406 Next, we assessed TILs by dissecting a single target liver metastatic lesion and a mixture of non-target liver metastases, respectively, in this model treated by IM injection of STING-A. 407 408 The single target liver lesion was pre-selected as it would be selected for the IT treatment, but without any IT treatment to be given(Fig.5E-H;Fig.S4D-E). As shown in Fig.5E-F, IM 409 injection of STING-A alone significantly enhanced the infiltration of CD8⁺ and CD4⁺T cells 410 in both the pre-selected liver metastatic lesion and other liver metastases. The enhancement 411 of the T cell infiltration was not as high in the IM combo group as the IM STING-A alone 412 group. Nevertheless, CD103⁺DCs showed a similar profile in the tumors with the IM 413

414 injection of STING-A(Fig.5G-H) compared to those shown above with the IT injection of
415 STING-A, suggesting that systemic IM injection of STING-A is able to activate the desired
416 antigen-presenting process in the liver metastases.

417

418 We then performed the NanoString analyses of liver metastases and SubQ tumors and 419 compared them among treatment groups. As shown in the Fig.S5A, the high-throughput 420 analysis of the NanoString results only revealed a smaller number of genes that were differentially expressed among the treatment groups than those differentially expressed 421 422 among the treatment groups with the IT injection of STING-A(Fig.S1E). The results however supported a non-significant increasing trend in the expression of genes associated with the 423 IFN response pathways in the liver metastases in the IM combo group(Fig.S5B). 424 425 Nevertheless, in the STING-A IM alone treatment group, the expression of T cell costimulatory molecules including Tnfrsf4(OX40), Cd27, Tnfrsf9(CD137), and Icos exhibited a 426 trend(Fig.S5C) consistent with the results in the single STING-A IT treatment group(Fig.3E). 427 However, the IM injection of STING-A combined with a-PD-1 did not significantly increase 428 the expression of these co-stimulatory molecules, together with the above T cell infiltration 429 results(Fig.5E-F), suggesting a-PD-1 or a-PD-1 alone may not be an optimal immune 430 checkpoint inhibitor treatment strategy in combination with STING-A. Nevertheless, we 431 observed a strong trend of increased expression of genes related to the cytotoxic activities of 432 effector T cells in the IM combo group including Gzma, Gzmb, Gzmk, and Prf1(Fig.S5D). 433 Chemokines especially those involved in effector T cell trafficking including Cxcl9, Cxcl10, 434 and Cxcl11 exhibited similar trend as those genes associated with cytotoxic activities of 435

effector T cells(Fig.S5E). In contrast, the increase in the T cell exhaustion and immune
checkpoint signals and myeloid cell-recruiting cytokine/chemokine signals that were
observed with the IT combo treatment were not observed in the IM combo group(Fig.6A).

439

440 We also examined the immune response in the SubQ tumors in the experiment with the IM 441 injection of STING-A. As shown in Fig.S6A, we observed a significant decrease in the infiltration ratio of CD4⁺PD-1⁺T cells in the CD4⁺T cells in the SubQ tumors from the IM 442 combo treatment group. A similar decreasing trend of CD8⁺PD-1⁺T cells was 443 444 observed(Fig.S6A), demonstrating the treatment effect of a-PD-1. Next, we used the same NanoString assays to assess the T cell functional status within the SubQ tumors. Interestingly, 445 unlike the results in the liver metastases, the expression of genes associated with the T-cell 446 447 receptor CD3 complex exhibited a significant increase (Cd3z) or a strong increasing trend in the SubQ tumors from the combo treatment group compared to other treatment 448 groups(Fig.S6B). T cell co-stimulatory molecules including *Tnfrsf4*(OX40), *Tnfrsf9*(CD137), 449 Cd80, and Icos all exhibited an increasing trend(Fig.S6C). In addition, the IM combo 450 treatment group exhibited a significantly increased expression of *ll7r*(Fig.S6D), which has 451 452 been shown to play a critical role in the development of lymphocytes in the process known as V(D)J recombination[33]. These results suggest that there may be an intertumoral 453 heterogeneity in the immune response to the systemic administration of STING-A. However, 454 we found that the T cell exhaustion and immune checkpoint signals exhibited an enhanced 455 456 expression in the IM combo treatment group compared to other treatment groups(Fig.6B). These results suggest that an enhanced T cell activation status and cytotoxic function in 457

response to either IT or IM treatment of STING-A is associated with upregulation of T cellexhaustion signals and CTLA-4.

460

The overall increasing trend of those chemokines/chemokine receptors involved in the 461 462 myeloid cell recruitment was less significant in both liver metastases(Fig.S5E) and SubQ tumors(Fig.S6D) than that in the IT combo group, suggesting that systemic delivery of 463 STING-A does not lead to a strong induction of immunosuppressive signals. It is noteworthy 464 to mention that Ccr5 expression in the SubQ tumors was significantly reduced by the 465 466 treatment of a-PD-1 compared to the vehicle treatment group, but was significantly enhanced following the IM injection of STING-A in combination with a-PD-1(Fig.S6E). This result 467 appears to be in consistent with the agonistic effect of C-C motif chemokine receptor(CCR) 5 468 469 expression previously reported in the PDAC models[34].

470

In addition to above differences between tumors from IM treated groups and IT treated 471 472 groups, we noted that the expression of immunosuppressive myeloid cells associated-genes including Apoe, Trem2, Cxcr4, and Abcg1 was significantly increased in the SubQ tumors 473 474 from the IM combo treatment group, but not the SubQ tumors from the IT combo treatment group, compared to the respective control group(Fig.6C-F). Such a difference was not 475 observed in the comparison between liver metastases from the IM combo treatment group 476 and the IT combo treatment group(Fig.S6F-I). These results suggest that systemic 477 administration of STING-A could still induce certain immunosuppressive signals that require 478 additional targeted treatments. 479

481 STING-A in combination with immune checkpoint inhibitors significantly improved the
482 survival of genetically engineered KPC mice that develops invasive PDAC
483 spontaneously.

484 Thus, above results suggest that systemic IM administration of STING-A led to less induction 485 of immunosuppressive signals while maintaining a systemic and intratumoral immune response that favors antitumor response. These results also provide the mechanistic basis of 486 the enhanced anti-tumor activity of the IM STING-A observed in the implanted tumor 487 488 model(Fig.5). We next validated this antitumor activity in the genetically engineered KPC mice that develops invasive PDAC spontaneously in a manner resembling human PDAC 489 pathogenesis. As CTLA-4 remains to be one of immunosuppressive signals presented in the 490 491 tumors treated by IM STING-A, we included anti-CTLA4 antibody to the immune checkpoint inhibitor regimen. KPC mice were subjected to weekly to twice weekly ultrasonic 492 screening at age of 3 months and enrolled in the experiment once either the length, width, or 493 height of the pancreatic tumor reached 2.00 mm to ensure eligible mice had equivalent tumor 494 burdens. As shown in Fig.7A, eligible KPC mice were randomized into the three treatment 495 496 groups to receive a total of seven injections. Treatment toxicity and mouse survival were monitored for 3 months following the first treatment. No treatment related toxicity including 497 local toxicity related to IM injection sites was observed. The STING-A monotherapy did not 498 499 show any antitumor activity in a later experiment(manuscript in preparation). Dual checkpoint inhibitors failed to improve the overall survival of KPC mice when compared 500 with control group; however, the co-administration of IM STING-A with dual checkpoint 501

inhibitors significantly prolonged the survival of the KPC mice(Fig.7B).

503

504 Systemic administration and intratumoral administration of an NLRP3 agonist in 505 combination with anti-PD-1 antibody has a similar efficacy

506 In this study, we attempted to understand whether systemic administration can be applied to another innate immune agonist. We found that the combination of IT injection of NACHT, 507 508 LRR, and PYD domains-containing protein 3(NLRP3) agonist and a-PD-1 had similar antitumor efficacy as the combination of IT injection of STING-A and a-PD-1 and prolonged the 509 510 survival of liver metastasis mice implanted with SubQ tumors as compared to a-PD-1 alone, NLRP3 agonist alone, or vehicle control(Fig.S7A). Likely due to the small sample size, the 511 combination of NLRP3 agonist and a-PD-1 failed to induce a significantly stronger tumor 512 513 growth inhibition on either liver metastatic lesion or the SubQ tumors than single treatment groups(Fig.S7B-C). However, different routes of administration of NLRP3 agonist including 514 IT, tail-vein injection, or intra-subcutaneous tumor injection resulted in no survival difference 515 516 of the liver metastasis mice(Fig.S7D). Despite a small sample size, NanoString analysis showed that the expression of Gzmk(Fig.S7E), an effector T cell cytotoxicity-associated gene, 517 was significantly increased in the NLRP3 agonist-treated tumor, but not in the STING-A-518 treated tumor(Fig.3D), suggesting that further investigation of systemic administration of 519 other innate immune agonists such as NLRP3 agonist is warranted. 520

521

522 Discussion

523 To our knowledge, this study is one of the few exploring the effects of the STING-A in

524 combination with ICIs for the treatment of PDAC in a pancreatic liver metastasis murine model. We previously used this mouse model to support the application of STING-A as an 525 526 adjuvant for the vaccine therapy. This mouse model resembles the spontaneous development of liver metastases in human PDACs and has been used in multiple prior studies for the 527 528 preclinical development of rationale immunotherapy combinations[35, 36, 37]. The TME of 529 the liver metastases in this model is similar to that of orthotopically implanted KPC 530 tumors[38]. Our study demonstrated the antitumor efficacy of the combinational treatment with STING-A and PD-1 blockade in this liver metastasis model as well as novel evidence 531 532 that demonstrated the survival benefit of the combination of a STING-A and ICIs in the genetically engineered, spontaneously formed KPC tumor model. In our model, STING-A in 533 combination with a-PD-1 enhances effector T cells infiltration and CD103⁺dendritic cells in 534 535 both target liver metastatic lesions and non-target metastatic lesions as well as distant tumor lesions that are resembled by subQ tumors. More importantly, this study demonstrated that 536 systemic administration of STING-A is equivalent to the intratumoral injection in both 537 antitumor efficacy and immune response. Neither systemic nor intratumoral administration 538 resulted in obvious toxicities; however, systemic administration was more feasible than 539 intratumoral administration and also avoided any intragenic liver injury. Our study also 540 supported the feasibility of administrating an NLRP3 agonist systemically and would 541 supports a new paradigm of the clinical development of innate immune agonists by systemic 542 administration. 543

544

545 Our results suggest that IM injection of STING-A is able to induce similar systemic immune

responses as the IT injection of STING-A. Systemic IM injection of STING-A is also able to activate the desired antigen-presenting process in the liver metastases. Although the effector T cell responses appear to be slightly weaker in the IM injection of STING-A, the increase in the T cell exhaustion and immune checkpoint signals and myeloid cell-recruiting cytokine/chemokine signals that were observed with the IT combo treatment were not observed in the IM combo group.

552

553 Although this study did not demonstrate a significant difference in the treatment response 554 between IM and IT injections of STING-A, we observed that mice who received IM injections of STING-A in combination with a-PD-1 survived longer than 6 weeks whereas 555 none of the mice who received IT injection of STING-A in combination with a-PD-1 556 557 survived longer than 6 weeks. Therefore, it might be possible to see the survival benefit of the IM injection of STING-A if the sample size would potentially be larger; however, the sample 558 size in each experiment was limited by the technical difficulty of IT injection. As it would be 559 a challenge to breed a large number of the KPC transgenic mice for being randomized to 560 multiple treatment groups, we decided to test the combination of a-PD-1 and anti-CTLA-4 561 562 antibody instead of two ICIs by itself in the experiments with the KPC transgenic mice. Our results suggest that an enhanced T cell activation status and cytotoxic function in response to 563 either IT or IM treatment of STING-A is associated with CTLA-4 upregulation. Hence, the 564 reason we included anti-CTLA-4 antibody to a-PD-1 in the experiment with the KPC 565 transgenic mice. Future studies could compare the combination of a-PD-1 and anti-CTLA-4 566 antibody with either a-PD-1 alone or anti-CTLA-4 antibody alone. 567

This study has a few limitations, first being the smaller sample size. As discussed above, the 569 sample size is limited by the technical complexity of IT injection and breeding KPC 570 transgenic mice. However, the sample size in the current study is, to our knowledge the 571 572 largest one in testing the intratumor injection of the tumors in internal organs. It is also one of 573 the largest using the KPC transgenic mice as the preclinical model for the efficacy testing. However, the current sample size allowed this study to repeat most of the experiments. 574 Second, this study did not examine the effect of anti-CTL4-4 antibody separately from the 575 576 treatment groups testing the combination of a-PD-1 and anti-CTLA-4 antibody. Third, chemotherapy, as a current standard of care treatment, was not included in the experiments in 577 this study. We were concerned that chemotherapy would complicate our comparison of the 578 579 antitumor efficacy and immune response between the IM and IT injection of STING-A. We are studying the combination of chemotherapy and STING-A in an independent study. 580

581

582 To test the feasibility of systemic administration of other innate immune agonists, we examined the NLRP3 agonist in this study. Although it may have been limited by the small 583 584 sample size, this study demonstrated that there were no significant differences in anti-tumor efficacy between intratumoral and systemic administration of NLRP3 agonist. This study has 585 not performed an in-depth investigation on the NLRP3 agonist. However, with limited data, 586 this study suggests that further investigation of systemic administration of other innate 587 immune agonists such as NLRP3 agonist is warranted. The results in this study have thus 588 supported the phase 1 trial that evaluated BMS-986301 intratumoral or intravenous injection 589

as monotherapy or in combination with nivolumab(PD-1 blockade) and ipilimumab inpatients with advanced solid cancers(NCT03956680).

592

Abbreviations: PDAC, pancreatic ductal adenocarcinoma; TME, tumor microenvironment; 593 594 STING, stimulator of interferon genes; PD-1, programmed cell death protein 1; CTLA-4, 595 cytotoxic T-lymphocyte associated protein 4; KPC, Kras/p53/pdx1-Cre; NLRP3, NACHT, LRR, and PYD domains-containing protein 3; TBK1, TANK-binding kinase 1; IRF3, 596 interferon regulatory factor 3; IFN, interferon; CDN, cyclic dinucleotide; ICI, immune 597 598 checkpoint inhibitors; TGI, tumor growth inhibition; TIL, tumor infiltrating leucocyte; DC, dendritic cell; IL, interleukin; CXCL, C-X-C motif chemokine ligand; CXCR, C-X-C motif 599 chemokine receptor; CCL, C-C motif chemokine ligand; G-CSF, granulocyte colony 600 601 stimulating factor; CCR, C-C motif chemokine receptor.

602

603 Declarations

604 Ethics approval and consent to participate

All studies and maintenance of mice were conducted in accordance with the approval of the
Institutional Animal Care and Use Committee (IACUC) guidelines of Johns Hopkins School
of Medicine (Animal Protocol: MO22M59).

608

609 **Consent for publication**

610 Not applicable.

612 Availability of data and materials

All data needed to evaluate the conclusions in the paper are present in the paper and the
Supplementary Materials. Any further information required to support our data will be
supplied upon request.

616

617 **Competing interests**

L.Z. receives grant support from Bristol-Meyer Squibb, Merck, AstraZeneca, iTeos, Amgen,
NovaRock, Inxmed, Halozyme and Abmeta. L.Z. is a paid consultant/Advisory Board
Member at Biosion, Alphamab, NovaRock, Ambrx, Akrevia/Xilio, QED, Novagenesis, Snow
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623

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632

633 Authors' contributions

634	Keyu Li and Junke Wang contributed equally to this project. Concept was conceived by L.Z.
635	The strategy and the overall study were designed by K.L. and L.Z. Experiments were
636	conducted by K.L., Junke W., N.N., N.J., A.B., Jiangxin W., M.H., J.Y., D.T., B.E., C.Y., and
637	M. W. Data were collected by K.L. and Junke W. Formal analysis was conducted by K.L.,
638	Junke W, and Y.X. Genetically modified mice were generated by M.H, V.F., and M.L.
639	Original draft manuscript was written by K.L. Manuscript was reviewed and revised was by
640	A.O., B.E., M.W., and L.Z. Supervision was made by X.L., Y.W., and L.Z. The project
641	administrator is L.Z.
642	
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647	
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742 Figures and Figure legends



745 Figure 1. Intratumoral injection of the STING agonist in combination with anti-PD-1 antibody enhance local anti-tumor efficacy with abscopal effects. (A) Ultrasonographic 746 747 measurement for the target liver metastatic lesion before and after intratumoral injection of the STING agonist. Arrow indicates the injection needle path. (B) Treatment schema. Mice 748 that met inclusion criteria were randomly assigned to each group on day 12 and monitored 749 twice a week until survival endpoints were met. (C) Tumor Growth Inhibition (TGI) of the 750 injected liver metastatic lesion during the treatment period. (D) Tumor Growth Inhibition of 751 the remote subcutaneous (SubQ) tumor during the treatment period. Dashed line at -50% 752 indicates statistically significant TGI. (E) Kaplan-Meier survival curves compare different 753 754 treatment groups. NC, vehicle/isotype antibody control; STING A, STING agonist; a-PD-1 755 Ab, anti-PD-1 antibody; Combo, STING A+a-PD-1 Ab; IT, intratumoral; IM, intramuscular; 756 IP, intra-peritoneal. Data shown as mean \pm SD; comparison by unpaired t test in C and D, and 757 by Log-rank test in E; p < 0.05; p < 0.01; NS, not significant. 758



Figure 2. STING agonist in combination with anti-PD-1 antibody enhances effector T cells 761 and CD103⁺dendritic cell (DC) infiltration and activates anti-tumor immunity via innate 762 immune response signaling pathways. (A) Percentages of the CD8⁺ and CD8⁺PD-1⁺T cells 763 among CD45⁺leucocytes in the target liver metastatic lesion. (B) Percentages of the 764 MHCII⁺CD11c⁺DC among CD45⁺leucocytes and the percent of CD11b⁻CD103⁺DC subtype 765 766 in the target liver metastatic lesion. (C) Percentages of the CD8⁺, CD4⁺, and CD4⁺PD-1⁺T 767 cells among CD45⁺leucocytes in the non-target liver metastases. (D) Percentages of MHCII⁺CD11c⁺DC among CD45⁺leucocytes and the percent of CD11b⁻CD103⁺DC subtype 768 in the non-target liver metastases. (E) Expression of genes in the IFN-response pathways in 769 770 the target liver metastatic lesions from different treatment groups. (F) Expression of genes in the TNF-response pathways in the target liver metastatic lesions from different treatment 771 772 groups. (G) Expression of genes in the innate immune response pathways in the target liver metastatic lesions from different treatment groups. Data are shown as the mean ± SD; 773 comparison by unpaired t test; *p < 0.05; **p < 0.01; ***p < 0.001; ***p < 0.001. 774 775 Remaining comparisons are non-significant.



777 778

Figure 3. Differentially expressed genes in the target liver metastatic lesions between treatment groups. Expression of genes, as indicated, in the gene families of cytokines (A-B), chemokines (C), the activation of effector T cells (D), the T cell co-stimulatory factors (E), and the immune checkpoint activators (F). Data shown as mean \pm SD; comparison by unpaired t test; *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001. Remaining comparisons are non-significant.



Figure 4. STING agonist in combination with anti-PD-1 antibody enhances the infiltration 788 and activation of effector T cells in the distant subcutaneous tumors. (A) Percentages of 789 immune cells in the remote SubQ tumors, such as the CD8⁺, CD8⁺PD-1⁺, CD4⁺, and 790 CD4⁺PD-1⁺T cells among CD45⁺leucocytes, the CD8⁺PD-1⁺among CD8⁺T cells, and the 791 CD4⁺PD-1⁺T cells among CD4⁺T cells, respectively. (B) Percentages of the 792 793 MHCII⁺CD11c⁺DC, CD11b⁻CD103⁺ and CD11b⁺CD103⁻ DC subtypes DC among CD45⁺leucocytes, respectively, in the distant SubQ tumors. Differentially expressed genes in 794 the distant SubQ tumors from different treatment groups, including those in the gene families 795 of cytotoxic T cell function (C), T cell activation (D), and the immune checkpoint activators 796 (E). (F) Heatmap of the interleukin family genes and chemokine genes that were significantly 797 increased in the Combo group comparing to the vehicle control treatment group. (G) 798 799 Expression of genes in the IFN-response pathways in SubQ tumors from different treatment groups. (H) Expression of genes in the TNF-response pathways in SubQ tumors from 800 different treatment groups. (I) Expression of genes in the innate immune response pathways 801 in SubQ tumors from different treatment groups. Data shown as mean \pm SD; comparison by 802 unpaired t test; *p < 0.05; **p < 0.01; ***p < 0.001; ***p < 0.001. Remaining 803 comparisons are non-significant. 804



808 Figure 5. Intramuscular injection of STING agonist in combination with anti-PD-1 antibody prolonged the survival of liver metastasis mice and induces both systemic and intratumoral 809 810 immune response. (A) Kaplan-Meier's survival curves compare the survival in different 811 intratumoral (IM) treatment groups. (B) TGI on distant SubQ tumors. Dashed line at -50% indicates statistically significant inhibition. (C) Comparison of serum concentrations of TNF-812 α and IFN- γ collected 6 hours after the first IM injection between treatment groups. (D) 813 Comparison of serum concentrations of CXCL9, CXCL10, IL-2, and IL-12 collected 6 hours 814 after the first IM injection between treatment groups. Percentages of the CD8⁺, CD8⁺PD-1⁺, 815 CD4⁺, and CD4⁺PD-1⁺T cells among CD45⁺leucocytes in the pre-selected, target single liver 816 817 metastatic lesions (E) and non-target liver metastases (F). Percentages of the MHCII⁺CD11c⁺DC and CD11b⁻CD103⁺subtype among CD45⁺leucocytes, respectively, in 818 the pre-selected single liver metastatic lesion (G) and non-target liver metastases (H). Data 819 shown as mean ± SD; comparison by Log-rank test for A and by unpaired t test for others; *p 820 < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001. Remaining comparisons are non-821 significant. 822





Figure 6. Comparison of expressions of differentially expressed genes in tumors from systemically treated mice. Expression of genes of the checkpoint gene family in pre-selected, target single liver metastatic lesions (A) and distant subcutaneous tumors (B) from the intramuscularly treated mice. Expression of *Apoe* (C), *Trem2* (D), *Cxcr4* (E), and *Abcg1* (F) in the distant subcutaneous tumors from both intratumoral and intramuscularly treated mice. Data shown as mean \pm SD; comparison by unpaired t test; *p < 0.05; **p < 0.01; ***p < 0.001. Remaining comparisons are non-significant.





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Figure 7. STING agonist in combination with immune checkpoint inhibitors significantly 835 improved the survival of genetically engineered KPC mice. (A) Treat schema. (B) Kaplan-836 Meier survival curves compare overall survival between different treatment groups. NC, 837 838 vehicle/isotype antibody control; STING A, STING agonist; a-PD-1 Ab, anti-PD-1 antibody;

a-CTLA-4 Ab, anti-CTLA-4 antibody; IM, intramuscular; IP, intra-peritoneal. Data shown as 839

mean \pm SD; comparison by Log-rank test; *p < 0.05; **p < 0.01; NS, not significant. 840