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CD122-targeted interleukin-2 and αPD-L1 treat bladder cancer and melanoma via distinct mechanisms, including CD122-driven natural killer cell maturation

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ABSTRACT

Bladder cancer (BC) and melanoma are amenable to immune checkpoint blockade (ICB) therapy, yet most patients with advanced/metastatic disease do not respond. CD122-targeted interleukin (IL)-2 can improve ICB efficacy, but mechanisms are unclear. We tested aPD-L1 and CD122-directed immunotherapy with IL-2/alL-2 complexes (IL-2c) in primary and metastatic bladder and melanoma tumors. IL-2c treatment of orthotopic MB49 and MBT-2 BC generated NK cell antitumor immunity through enhanced activation, reduced exhaustion, and promotion of a mature, effector NK cell phenotype. By comparison, subcutaneous B16-F10 melanoma, which is IL-2c sensitive, requires CD8⁺ T and not NK cells, yet we found α PD-L1 efficacy requires both CD8⁺ T and NK cells. We then explored αPD-L1 and IL-2c mechanisms at distinct metastatic sites and found intraperitoneal B16-F10 metastases were sensitive to aPD-L1 and IL-2c, with IL-2c but not aPD-L1, increasing CD122⁺ mature NK cell function, confirming conserved IL-2c effects in distinct cancer types and anatomic compartments. αPD-L1 failed to control tumor growth and prolong survival in B16-F10 lung metastases, yet IL-2c treated B16-F10 lung metastases effectively even in T cell and adaptive immunity deficient mice, which was abrogated by NK cell depletion in wild-type mice. Flow cytometric analyses of NK cells in B16-F10 lung metastases suggest that IL-2c directly boosts NK cell activation and effector function. Thus, aPD-L1 and IL-2c mediate nonredundant, immune microenvironment-specific treatment mechanisms involving CD8⁺T and NK cells in primary and metastatic BC and melanoma. Mechanistic differences suggest effective treatment combinations including in other tumors or sites, warranting further studies.

Background

Bladder cancer (BC) and melanoma are among the most common cancers in the United States, collectively accounting for an estimated 190,000 new cancer diagnoses and 25,000 deaths annually.¹ Nearly all cancer-related mortality is from metastases^{2–4} and 5-year survival rates for patients with metastatic BC and melanoma are poor at 6% and 30%, respectively.⁵ BC and melanoma share common metastatic sites, including regional lymph nodes, bones, and lung.^{6,7} Melanoma is also one of the most common extra-abdominal malignancies to develop intraperitoneal metastasis, which is uniformly associated with a poor prognosis.⁸

Immune checkpoint blockade (ICB) antibodies targeting programmed death receptor 1 (PD-1) or one of its ligands (programmed death ligand 1, PD-L1), and cytotoxic T-lymphocyte-associate protein 4 (CTLA-4) have provided durable clinical efficacy in patients with advanced or metastatic melanoma, yet more than 50% do not respond to ICB.^{9–12} BC, like melanoma, is thought to be among the most ICB-responsive cancers due to high tumor mutational burden, yet only 10%–30% of BC patients respond to ICB.¹³

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High dose interleukin-2 (IL-2) became a United States Food and Drug Administration-approved metastatic melanoma immunotherapy in 1998, yet is highly toxic with low overall efficacy, in part because of its promotion of regulatory T cells (Tregs).^{14,15} IL-2 signals by binding to the IL-2 receptor (IL-2R), which exists as either a trimer of IL-2R α (CD25), IL-2R β (CD122), and IL-2 R γ (γ c, CD132) subunits or as a dimeric $\beta\gamma$ IL-2 R.^{16,17} Activated T cells and CD4⁺ Foxp3⁺ Tregs express trimeric receptors,¹⁸ whereas memory-phenotype CD8⁺ CD44⁺ T cells and NK cells highly express dimeric $\beta\gamma$ IL-2 Rs and can mediate anti-tumor immunity.¹⁷ Selective dimeric $\beta\gamma$ IL-2R stimulation with CD122-directed IL-2/ α IL-2 complexes (IL-2c) boosts antitumor immunity in murine cancer models by augmenting CD122⁺ effector T cell functions while avoiding the deleterious Treg-promoting effects of IL-2 monotherapy.¹⁹

However, specific effector populations activated by IL-2c are highly dependent on tumor type and the tumor immune microenvironment, which also differentially affects α PD-L1 efficacy despite potentially targeting similar immune populations.²⁰ Here, we studied IL-2c and α PD-L1 immunotherapy treatment mechanisms in orthotopic mouse models

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Supplemental data for this article can be accessed on the publisher's website

of BC (MB49 and MBT-2) and melanoma (B16-F10) versus mechanisms in peritoneal and lung metastatic sites. We show that IL-2c promotes NK cell maturation and effector function in distinct primary (orthotopic bladder) and metastatic (peritoneum and lung) sites in a T cell-independent manner. NK cells were indispensable for efficacious treatment of lung metastatic BC and melanoma. Notably, IL-2c effects on CD8⁺ T cells are complementary with aPD-L1 in metastatic melanoma models and treatment effects show tumor and anatomic compartment-dependence. These data shed light on conserved mechanisms of selective IL-2 receptor targeting on NK cell functions and help define treatment response mechanisms, in addition to potentially responsive patient populations, as CD122-directed immunotherapy and ICB combinations are currently in human trials.²¹

Methods

Mice

Wild-type C57BL/6J (BL6) and C3H/HeJ (C3H) mice and B6.129P2-Tcrd^{tm1Mom}/J (TCR δ^{KO}), B6.129P2-Tcrb^{tm1Mom}/J (TCR β^{KO}), and B6.129S7-Rag1^{tm1Mom}/J (RAG1^{KO}) genetic knock out mice all on the BL6 background were purchased from The Jackson Laboratory (Bar Harbor, ME). Mice were bred in our animal facility, given *ad libitum* water and food, and housed under specific pathogen-free conditions. All mice were at least 8 weeks old and age- and sex-matched when used for each experiment.

Tumor cell lines and cell culture

Mouse MB49 and MBT-2 BC cell lines were engineered to express luciferase as we described.¹⁹ Mouse B16-F10 (herein B16) was purchased from ATCC (Manassas, VA). We developed Luciferase-expressing B16 using CMV-Firefly luciferase-IRES-Puro lentivirus from Cellomics Technology (Halethrope, MD), following the manufacturer's protocol. All cell lines are PD-L1⁺²²⁻²⁴ and were passaged <5 times prior to tumor challenge. Cells were maintained in 5% fetal bovine serum (FBS)-containing DMEM (Dulbecco's Modified Eagle Medium, MB49, and B16) or RPMI-1640 (Roswell Park Memorial Institute, MBT-2), supplemented with 1/100 dilutions of penicillin/streptomycin, L-glutamate, and 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES).

In vivo tumor challenges

Orthotopic (intravesical) BC challenge with MB49 and MBT-2 used 8×10^4 or 1×10^6 cells, respectively, in 50 µL Dulbecco's phosphate buffered saline (DPBS, Sigma Aldrich), via indwelling urinary catheter in female mice under isoflurane anesthesia, as described.²⁵ Orthotopic (subcutaneous) B16 challenge used 5×10^5 cells in 200 µL DPBS into each flank of male mice (2 tumors/mouse). Subcutaneous B16 tumor growth was measured every 2 days by calipers and volume was calculated as (length x width²)/2. Intraperitoneal B16 challenge used 4×10^5

cells injected into the peritoneum. Lung BC tumors were generated via intravenous tail vein injection of 7×10^5 MB49 or 2.5×10^5 MBT-2 cells in 200 µL DPBS into male mice.²⁰ Lung B16 tumors were generated via similar methods using 3×10^5 B16 cells. Bladder tumor and peritoneal B16 tumor growth was assessed by tumor weight at sacrifice. Lung B16 tumor growth was measured *in vivo* by tumor bioluminescence every 4 days, starting on day 7 post challenge. Survival was defined as spontaneous death, moribundity, tumor volume $\geq 2000 \text{ mm}^3$ (subcutaneous B16), or significant weight loss from baseline ($\geq 15\%$ for bladder and $\geq 20\%$ for lung tumors). To ensure uniform tumor size across all treatment groups, mice were occasionally excluded upon first tumor measurement before randomization, if determined to be an outlier by Grubbs' test.

In vivo bioluminescence monitoring

Bioluminescent imaging (IVIS Lumina Imaging System; Perkin Elmer; Waltham, MA) was used for treatment randomization and *in vivo* tumor monitoring as described.¹⁹ In brief, mice were imaged 15 minutes after intraperitoneal injection of 3 mg PBS-dissolved d-luciferin K⁺ (Gold Biotechnology; St. Louis, MO) with one-minute exposure, small binning, and F/stop = 1. Identical regions of interest were drawn over each mouse and average radiance (photons/sec/cm²/sr) was quantified using Living Image software version 4.2.

In vivo treatments

αPD-L1 (clone 10 F.9G2), αIL-2 (clone JES6-5H4), and isotype control antibodies (clone LTF-2 rat IgG2b and polyclonal Armenian hamster IgG) were purchased from BioXCell (Lebanon, NH). Carrier-free recombinant mouse IL-2 was purchased from Biolegend (San Diego, CA). IL-2c is 1.5 µg/ mouse IL-2 complexed with 7.5 µg/mouse αIL-2 at a 1:2 molar ratio defined as optimal in PBS at 37°C for 15–30 min,²⁶ before intraperitoneal administration in 100 µL DPBS. IL-2c treatment was every other day for four doses. Treatment began on day 6 or 7 post challenge for orthotopic BC, subcutaneous melanoma, or peritoneal melanoma challenge, and day 7 or 8 for intravenous lung tumor challenge, unless noted otherwise. 100 µg/mouse αPD-L1 was given intraperitoneally every five days for three doses starting on day 7–9 post tumor challenge, with all experiments using relevant isotype controls.

In vivo cell depletions

α-asialo ganglioside GM1 (asGM1, clone Rabbit) was purchased from FUJIFILM Wako Chemicals U.S.A. Corporation (Richmond, VA) and prepared according to manufacturer's instructions using 1 mL of deionized water. αNK1.1 (clone PK136), αCD8 (clone 2.43), and isotype control antibodies were purchased from BioXCell. Antibodies were given intraperitoneally to deplete relevant cells concurrently with the tumor treatment regimen per figure legends. Doses/mouse: 200 μL α-asGM1, 250 μg αNK1.1, and 250 μg αCD8.

Flow cytometry

Tumors were harvested for flow cytometry after mice were sacrificed via cervical dislocation and induction of deep iso-flurane anesthesia. Bladder tumor dissection was as previously described.²⁰ Subcutaneous and peritoneal B16 tumors were dissected and placed in a 6-well plate filled with serum-free RPMI-1640 and manually dissociated with the back of a syringe. Cells were incubated for 45–60 min in 3 mL serum-free RPMI-1640 with 0.25 mg/mL DNAse I and 1.65 mg/mL collagenase Type IV (both Sigma Aldrich; St. Louis, MO) and passed through a 70 µm filter to generate single-cell suspensions. $3-5 \times 10^6$ cells were then transferred to 96-well plates and samples with $<5 \times 10^6$ cells from the same group were pooled to ensure uniform cell counts for all samples.

Single-cell suspension from lung was prepared as previously reported.²⁷ Briefly, lungs bearing B16 tumors were excised from mice and lung tissue was dissociated via razor blades into fragments of ~1 mm in size. Then, tissue fragments were transferred into a 24-well plate containing 2 mL/lung of digestion medium (complete RPMI containing 2 mg/mL collagenase IV + 0.02 mg/mL DNase I) and incubated at 37°C for 30 min. Cells were then passed through a 70 μ m filter and a single-cell suspension was generated similar to subcutaneous and peritoneal B16 tumors.

Dead cells were excluded using either LIVE/DEAD Fixable Blue Dead Cell Stain Kit for ultraviolet excitation (Thermo Fisher Scientific; Waltham, Massachusetts, USA) or Ghost Dye Violet 510 (Tonbo biosciences; San Diego, California, USA). Fc blocking to prevent nonspecific labeling was done using 1:100 dilution anti-CD16/32 (clone 93, Biolegend). A list of used antibodies, including surface and intracellular staining, is provided in Table 1. Cells were stained for surface antigens by incubating at 4°C for 30–45 min with antibodies against: CD45 (30-F11), CD3 (17A2/145-2C11), CD8 (53–6.7), B220 (RA3-6B2), PD-1 (J43), CD27 (LG.3A10), CD11b (M1/70), CD69 (H1.2F3), CD49b (HM α 2), CD122 (TM- β 1), KLRG1 (2F1), NK1.1 (PK136), DNAM-1 (10E5), NKG2A (16a11), and gp100 [EP4863(2)], all at manufacturer recommended dilutions.

For intracellular staining, cells were fixed and permeabilized with FoxP3/transcription factor buffer (eBioscience; San Diego, CA) according to manufacturer instructions, and incubated at 4°C for 45 min. To assess effector molecule production, cells were stimulated with Cell Activator Cocktail (Biolegend) containing phorbol 12-myristate 13-acetate, ionomycin, and brefeldin A at 2 µL cocktail/mL CR10 medium (RPMI-1640 with 10% FBS, L-glutamine, sodium pyruvate, non-essential amino acids, penicillin/streptomycin, and HEPES buffer) for 6 hours in a 37°C incubator. Following stimulation, intracellular staining was performed by incubating cells at 4°C for 30-45 min with antibodies against: Eomes (W17001A), Perforin (S16009A), Granzyme B (NGZB/GB11), Interferon (IFN)-y (XMG1.2), all at manufacturer recommended dilutions in FoxP3 permeabilization buffer. Brilliant Stain Buffer Plus (BD) was added to surface and intracellular staining at the manufacturer recommended dilution. Absolute cell numbers were determined by multiplying the cell ratio per live, singlet cell in each flow sample by total cell numbers in the sample.

			,	, ,	
			Catalog		
Antibodies	Species	Clone	Fluorochrome	number	Manufacturer
CD45	Mouse	30-F11	AF532	58-0459-	eBioscience
		20 511		42	Diala waad
	Mouse	30-F11	APC/FIRE 810	103174	Biolegena
600	Mouse	30-F11	AF/00	103128	Biolegend
CD3	Mouse	1/A2	BV510	100233	Biolegend
	Mouse	145-	PE-Cy5	55-0031-	lonbo
		2 C11		U100	bioscience
CD8a	Mouse	53–6.7	BUV395	563786	BD
	Mouse	53–6.7	APC-Cy7	25–0081-	Tonbo
				U025	bioscience
B220	Mouse	RA3-6B2	BUV661	612972	Biolegend
PD-1	Mouse	J43	PerCP-	46–9985-	eBioscience
			eFlour710	82	
	Mouse	J43	BB700	566515	BD
CD27	Mouse/rat/	LG.3A10	BV605	124249	Biolegend
	human				-
CD11b	Mouse	M1/70	AF700	101222	Biolegend
	Mouse/	M1/70	BV570	101233	Biolegend
	human				je v sje v
CD69	Mouse	H1.2F3	PF/Dazzle 594	104535	Biolegend
	Mouse	H1.2F3	BV650	104541	Biolegend
	Mouse	H1 2F3	BUV563	741234	BD
CD49b	Mouse	HMa2	BV711	740704	BD
CD122	Mouse	TM-B1	BV786	740869	BD
CD122	Mouse	TM-B1	DF DF	123210	Biolegend
	Mouse			123210	Biologond
	Mouse	7E1	PVAD1	566294	PD
	Mouse			741026	
	Mouse	PKIDO		100740	DU Dialamand
	Mouse	PK130	BV/85	108749	Biolegena
	Mouse	PK136	APC	20-5941-	Ionbo
60 a a 4			DE /D E0.4	0100	bioscience
CD226 (DNAM-	Mouse	10E5	PE/Dazzle 594	128818	Biolegend
	Mouro	16-11	DorCD	46 5907	Pieccionco
INKGZABO	Mouse	10011	eFlour710	40-5897- 82	ebioscience
gp100	Mouse	EP4863 (2)	AF647	ab246730	abcam
Fomes	Mouse	W17001A	AF647	157703	Biolegend
Perforin	Mouse	S16009A	PF	154306	Biolegend
1 chionini	Mouse	S16009A	APC	154304	Biolegend
Granzyme	Mouse/	GR11	Pacific Blue	515408	Biolegend
R	human	GUTT	rucine Diue	00701	Diolegena
D	Mouse	NG7R	PE-Cv5 5	35_8808	eRioscience
	MOUSE	NULD	1 L CYJ.J	82	CDIOSCICIICE
IFN-γ	Mouse	XMG1.2	BUV737	612769	BD

Flow data were acquired on a Cytek Aurora flow cytometer (Cytek Biosciences; Freemont, CA) and analyzed using FlowJo software (BD) version 10.7.1. Representative flow cytometry gating strategies and relevant immune phenotype comparisons across tumor models are shown in *Supplementary Data* and Table 2, respectively.

Statistical and data analyses

Data were analyzed and graphed with GraphPad Prism 9.1.2. Data with error bars are mean \pm SEM. To compare two means, we used an unpaired *t* test. Three or more means were compared with one-way ANOVA and *post hoc* Sidak's test. Tumor growth curves were compared by two-way ANOVA, analyzed for overall treatment effect, followed by *post hoc* Sidak's test of discrete time points. Log-rank test was used to compare Kaplan-Meier survival curves. Occasionally, data sets with suspected outliers were identified by Grubbs' test (used only

Table 2. Flow	cytometric assessment	of NK and CD8 ⁺ T	Γ cell phenotypes across	tumor models
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		Measurement						
Tumor	Cell			Inhibitory				Additional
model	type	Prevalence	Cell number	receptor	Activation	NK maturation	Function	Markers
MB49 WT	NK cell	% NK1.1 ⁺ of CD3 ⁻	N/A	PD-1/NKG2A	CD69/ DNAM- 1	% CD27 ⁻ of NK % CD69 ⁺ Eomes ⁺ CD27 ⁻ of NK	Perforin	N/A
MBT- 2 WT	NK cell	N/A	N/A	N/A	N/A	% CD27 ⁻ NK % KLRG1 ⁺ of CD27 ⁻ NK	N/A	N/A
MB49 TCRδ ^{KO}	NK cell	% NK1.1 ⁺ of CD3 ⁻	% x Live Cell Count	PD-1	N/A	% and # KLRG1 ⁺ of Eomes ⁺ CD27 ⁻ NK	Perforin	CD11b, CD49b
B16 SQ	CD8 ⁺ T cell	N/A	Normalized to tumor weight	N/A	N/A	N/A	Perforin	N/A
	NK cell	% NK1.1 ⁺ of CD3 ⁻	N/A	PD-1	N/A	N/A	Perforin	N/A
B16 IP	CD8 ⁺ T cell	% CD8 ⁺ of CD3 ⁺	N/A	N/A	N/A	N/A	Perforin/IFN-γ/ Granzyme B	CD122
	NK cell	N/A	Normalized to tumor weight	PD-1	CD69	% CD69 ⁺ Eomes ⁺ CD27 ⁻ of NK	Perforin/Granzyme B	CD122
B16 Lung	CD8 ⁺ T cell	% CD8 ⁺ of CD3 ⁺	Normalized to gp100 ⁺ cell number	N/A	CD69	N/A	IFN-γ/Granzyme B	N/A
	NK cell	% NK1.1 ⁺ of CD3 ⁻	Normalized to gp100 ⁺ cell number	N/A	CD69	% KLRG1 ⁺ of NK	IFN-γ/Granzyme B	CD122

WT, wild type. N/A, not applicable. SQ, subcutaneous. IP, intraperitoneal.

once for a given data set) and removed from analysis. For all analyses, significance was based on a multiplicity-corrected α of 0.05.

Results

IL-2c enhances NK cell activation and maturation in orthotopic BC

We previously reported that CD122-targeting IL-2c activated $\gamma\delta$ T cells but not CD8⁺ T cells to treat orthotopic MB49 BC.²⁰ However, effects on other CD122⁺ immune cell populations in bladder are unreported. Like MB49, we observed reduced tumor growth and tumor weight in IL-2c treated mice bearing orthotopic MBT-2 (Fig. S1A,B), a syngeneic mouse BC line on a distinct genetic background (C3H/He) from MB49 (BL6). To assess NK cell effects in IL-2c treatment, we performed flow cytometric analysis of bladders from mice bearing orthotopic MB49 and MBT-2 tumors (Fig. S2). Compared to isotype treatment, IL-2c increased intratumoral NK cell frequency (Figure 1a), reduced PD-1 expression (Figure 1b) and increased CD69 expression (Figure 1c), corresponding with reduced exhaustion and enhanced activation, respectively. We also observed a trend toward decreased expression of the NK cell inhibitory receptor (NKG2A) in IL-2c treated bladders, yet other NK activating receptors (e.g., DNAM-1) were unaffected by IL-2c treatment (Fig. S3A).

NK cell maturation describes the process of terminal differentiation and acquisition of optimal effector function and can be defined by the loss of CD27 expression in nonlymphoid tissues.²⁸ We noted a significant increase in the frequency of CD27⁻ NK cells (Figure 1d) in MB49 tumors, suggesting increased NK cell maturation after IL-2c treatment. Moreover, an increase in frequency of CD69⁺ Eomes⁺ CD27⁻ NK cells (Figure 1e) in the context of efficacious IL-2c treatment suggests that these activated, mature NK cells may be important mediators of the IL-2c driven antitumor immune response. We also observed a trend for increased perforin-producing NK cells after IL-2c treatment (Fig. S3B), further suggesting IL-2c driven improvement in NK cell effector function. Data in the MBT-2 model corroborated these results, with IL-2c also increasing CD27⁻ NK cell frequency (Fig. S3C) in this model. CD27⁻ NK cells in MBT-2 tumors also highly expressed KLRG1 (Fig. S3D), a marker only present in the final stage of NK cell maturation.²⁹ To test NK contributions to IL-2c efficacy directly, we treated wild-type mice with IL-2c \pm NK cell depletion using α -asialoGM1,³⁰ which significantly reduced treatment efficacy in orthotopic MB49 (figure 1f). Collectively, these data implicate NK cells as critical mediators of IL-2c antitumor efficacy in orthotopic bladder tumors through promotion of NK cell maturation and effector function.

IL-2c promotes NK cell maturation in a γδ T cellindependent manner

We previously reported that antibody-mediated or genetic depletion of yo T cells significantly abrogated IL-2c efficacy in orthotopic MB49 challenge,²⁰ similar to NK cell depletion shown here (figure 1f). Thus, we hypothesized that IL-2c treatment efficacy is either the summation of independent $\gamma\delta$ T and NK cell effects or the result of their coordinated interactions. To test interactions, we performed flow cytometric analysis of intratumoral bladder NK cell content in orthotopic MB49 challenged TCR δ^{KO} mice (lacking all $\gamma\delta$ T cells) ± IL-2c treatment. As in wild-type mice (Figure 1), IL-2c NK cell effects included increased intratumoral bladder NK cell frequency (Figure 2a), reduced PD-1 expression (Figure 2b), and increased frequency of perforin-producing NK cells (Figure 2c). We then evaluated if the NK cell maturation phenotype observed in IL-2c treated wild-type mice was also present in $TCR\delta^{KO}$ mice. As in wild-type mice, we found a robust increase in the frequency and number of NK cells co-expressing KLRG1 and Eomes (Figure 2d,e), both of which are essential for NK cell maturation and function.^{29,31} These data suggest that IL-2c driven NK cell maturation in this model appears to be $\gamma\delta$ T cell independent, suggesting potential benefits of IL-2c in other contexts.



Figure 1. IL-2c enhances NK activation and maturation to treat orthotopic BC. Wild-type female mice were challenged orthotopically with 8×10^4 MB49 and treated with IL-2c or an isotype control every other day starting on day 7 for a total of 4 doses. (a-e) Mice were sacrificed on day 16 post challenge. Flow cytometric analysis of intratumoral bladder NK cell frequency (a), exhaustion (PD-1⁺) (b), activation (CD69⁺) (c), maturation (CD27⁻) (d), and CD69⁺ Eomes⁺ expression by CD27⁻ NK cells (e). N = 6–8 pooled bladders/group. p, unpaired t test. NK cells in MB49 defined as CD45⁺ CD3⁻ NK1.1⁺ cells. (f) Mouse survival in orthotopic MB49 challenge treated with IL-2c (as above) \pm 200 µL α-asialo GM1 (asGM1). N = 7–9 mice/group. p, log-rank. Note: We previously reported data showing survival of isotype-treated mice bearing orthotopic MB49.²⁰ Eomes, Eomesodermin. IL-2c, interleukin-2 complex. NK, Natural Killer. PD-1, Programmed Death Receptor 1.

We then performed more detailed analyses of NK cell maturation in TCR δ^{KO} mice bearing orthotopic MB49 by examining CD49b⁺ NK cells, as CD49b is a marker acquired at the onset of NK cell maturation in BL6 mice.²⁹ Using differential expression of CD11b and CD27 to evaluate the final three stages of NK cell maturation,³² we found that IL-2c significantly increased the frequency of terminally differentiated, mature CD11b⁺ CD27⁻ NK cells (Fig. S4A,B), with striking increases in KLRG1⁺ NK cell frequency (Fig. S4C), as observed in wild-type hosts (Figure 1). IL-2c also increased the number of Perforin⁺ Eomes⁺ KLRG1⁺ mature NK cells (Fig. S4D) suggesting enhanced effector function.

aPD-L1 requires CD8⁺ T and NK cells to treat subcutaneous B16 melanoma

IL-2 activates downstream STAT5 signaling via CD122 in NK cells and facilitates NK cell cytotoxicity,³³ yet prior studies in subcutaneous B16 melanoma showed that CD8⁺ T cells but not NK cells, are required for IL-2c treatment efficacy.³⁴ To determine if lack of NK cell requirements for IL-2c treatment efficacy in orthotopic B16 is tissue microenvironment-specific, we studied aPD-L1 which has established effects on CD8⁺ T and NK cells in several cancers.³⁵ We confirmed aPD-L1 efficacy in B16 in our mice (Figure 3a), consistent with prior



Figure 2. IL-2c promotes NK cell maturation in a $\gamma\delta$ T cell independent manner. TCR δ^{KO} female mice were challenged orthotopically with 8 × 10⁴ MB49 cells and treated with IL-2c on days 9, 11, 13, and 15 or isotype control. Mice were sacrificed on day 17 for flow cytometric analysis of NK cell frequency (a), exhaustion (i.e., PD-1⁺) (b), perforin production (c), and KLRG1⁺ Eomes⁺ expression by CD27⁻ NK cells (d,e). NK cells in MB49 defined as CD45⁺ CD3⁻ NK1.1⁺ cells. N = 9–11 mice/group. Tumors were pooled to achieve 5 × 10⁶ live cells per sample. p, unpaired t test. Eomes, Eomesodermin. IL-2c, interleukin-2 complex. KLRG1, Killer Cell Lectin Like Receptor G1. NK, Natural Killer. PD-1, Programmed Death Receptor 1.

reports.^{36,37} Flow cytometric analysis of subcutaneous B16 tumors (Fig. S5) showed that aPD-L1 improved CD8⁺ T cell number (Figure 3b) and effector function detected as increased perforin expression (Figure 3c), consistent with enhanced T cell cytotoxicity. However, aPD-L1 also increased intratumoral NK cell frequency (Figure 3d), reduced PD-1 expression (Figure 3e), and increased prevalence of perforin-producing NK cells (figure 3f), showing the potential for beneficial aPD-L1 effects on NK cells in skin. We then depleted $CD8^+$ T cells with aCD8 (Fig. S6) or NK cells with a-asialoGM1 during aPD-L1 treatment and found that both depletions significantly abrogated aPD-L1 mediated tumor control of subcutaneous B16 melanoma (Figure 3g). Together, these data confirm distinct tumor microenvironment-dependent effects of IL-2c and aPD-L1 in treatment of orthotopic (primary) B16 and provide an important baseline for comparison with mechanisms of these agents at metastatic sites.

IL-2c or aPD-L1 alone treat peritoneal B16 melanoma

We have extensively studied orthotopic (peritoneal) ID8agg mouse ovarian cancer, which is refractory to α PD-L1 but sensitive to IL-2c.¹⁹ To compare IL-2c and α PD-L1 mechanisms in orthotopic (subcutaneous) B16 and ID8agg, we used a clinically relevant peritoneal metastatic melanoma model⁸ via intraperitoneal injection of B16 into wild-type mice.³⁸ Peritoneal tumors from α PD-L1 treated mice were marginally smaller than isotype controls, yet IL-2c treatment led to a dramatic reduction in tumor weight versus isotype and α PD-L1 treated mice (Fig. S7A). Both agents equally improved mouse survival (Fig. S7B).

IL-2c promotes NK cell maturation in peritoneal B16, whereas aPD-L1 does not

Flow cytometric analyses of peritoneal B16 tumors (Fig. S8) showed that IL-2c increased total intratumoral CD45⁺ immune cell content (Figure 4a) compared to isotype control and aPD-L1 treated mice. aPD-L1 and IL-2c shared many beneficial effects on CD8⁺ T cells including increasing their prevalence (Figure 4b) and production of effector molecules including perforin, IFN-y, and granzyme B (Figure 4c-e), yet IL-2c effects were superior to aPD-L1 in these regards (Figure 4b-e). IL-2c also had NK cell activating effects not elicited by aPD-L1, including increased NK cell concentration (Fig. S9A) and CD69 expression (Fig. S9B), though both agents reduced NK cell PD-1 expression versus isotype controls (Fig. S9C). IL-2c-mediated NK activation also increased effector molecule production (figure 4f,g), frequency of CD69⁺ Eomes⁺ CD27⁻ activated mature NK cells (Figure 4h), and increased NK cell CD122 expression (Figure 4i), effects not observed with aPD-L1 (figure 4f-i).

To assess CD8⁺ T and NK cell contributions to α PD-L1 and IL-2c treatment mechanisms, we administered α PD-L1 \pm CD8⁺ T and NK cell depleting antibodies to peritoneal B16 tumor-bearing



Figure 3. α PD-L1 requires NK and CD8⁺ T cells to treat subcutaneous B16 melanoma. Wild-type male mice were challenged subcutaneously with 5 × 10⁵ B16 cells on both flanks and treated with 100 µg α PD-L1 on days 7, 12, 17 or an isotype control (a-f). (a) Tumor growth of mice bearing subcutaneous B16. N = 7 mice/group. p, two-way ANOVA of day 13 tumor volume. (b-e) Mice were sacrificed on day 14 for flow cytometric analysis of CD8⁺ T cell number (b) and perforin production (c), and NK cell prevalence (d), exhaustion (i.e., PD-1⁺) (e), and perforin production (f). N = 6–7 mice/group. p, unpaired t test. (g) Wild-type male mice were challenged subcutaneously with 5 × 10⁵ B16 cells on a single flank and treated with 100 µg α PD-L1 on days 7, 10, and 13 ± 200 µL α-asGM1, 250 µg α CD8, or an isotype control every three days starting the day before the challenge until the end of α PD-L1 treatment. N = 5 mice/group. p, two-way ANOVA of day 17 tumor volume. ANOVA, analysis of variance. asGM1, asialo GM1. NK, Natural Killer. PD-L1, Programmed Death Ligand 1. PD-1, Programmed Death Receptor 1.

mice. Consistent with our flow cytometric analyses that showed α PD-L1 effects primarily via CD8⁺ T cells and not NK cells in this model, CD8⁺ T cell depletion significantly reduced survival of mice treated with α PD-L1, whereas NK cell depletion did not impact survival (Figure 4j). As CD8⁺ T cell depletion did not

fully abrogate treatment efficacy, other immune cells likely also contribute. IL-2c treatment of peritoneal B16 had similar CD8⁺ T cell requirements, yet surprisingly, NK cell depletion did not significantly reduce survival (Figure 4k). This may be due to the strength of IL-2c CD8⁺ T cell effects in this context compared to



Figure 4. IL-2c activates of CD8⁺ T cells and promotes NK cell maturation in peritoneal B16. Wild-type male mice were challenged peritoneally with 4×10^5 B16 cells and treated with 100 µg α PD-L1 on days 7, 12, 17 or IL-2c on days 7, 9, 11, 13. Mice were sacrificed on day 15 for flow cytometric analysis of immune (CD45⁺) cell number normalized to tumor weight (a). (b-e) Intratumoral CD8⁺ T cell frequency (b) and effector molecule production (c-e). (f-i) Intratumoral NK cell perforin (f) and granzyme B

(g) production, CD69⁺ Eomes⁺ expression by CD27⁻ NK cells (h), and CD122 expression (i). N = 11–13 mice/group. *p* value, one-way ANOVA. (j) Survival of wild-type male mice challenged peritoneally with 4×10^5 B16 cells and treated with α PD-L1 (as above) \pm 250 ug α NK1.1 or 250 ug α CD8 on days 6, 9, 12, 15, 18. N = 8–14 mice/group. *p* value, log-rank. (k) Survival of wild-type male mice challenged peritoneally with 4×10^5 B16 cells and treated with α PD-L1 (as above) \pm 250 ug α NK1.1 or 250 ug α CD8 on days 6, 9, 12, 15, 18. N = 8–14 mice/group. *p* value, log-rank. (k) Survival of wild-type male mice challenged peritoneally with 4×10^5 B16 cells and treated with IL-2c (as above) \pm 250 ug α NK1.1 or 250 ug α CD8 on days 6, 9, 12, 15. N = 8–9 mice/group. *p* value, log-rank. Note: Survival of mice in Figure 4j,k can be compared to survival of isotype-treated mice with peritoneal B16 in Fig. S7B. ANOVA, analysis of variance. Eomes, Eomesodermin. IFN- γ , interferon-gamma. IL-2, interleukin-2 complex. MFI, Mean Fluorescence Intensity. NK, Natural Killer. PD-L1, Programmed Death Ligand 1.

NK cells, as $CD8^+$ T cells expressed higher levels of CD122 (the IL-2c target) both at baseline and during IL-2c treatment (Fig. S9D). These data show that in addition to many shared effects on $CD8^+$ T cells with aPD-L1, IL-2c-mediated promotion of NK cell maturation and function extends across tumor types and anatomic locations, though specific contributions of pleiotropic IL-2c immune effects to immunotherapy efficacy is tumor microenvironment-specific.

IL-2c treats B16 lung metastases, whereas aPD-L1 is ineffective

The lung is the most common site of visceral metastasis in melanoma³⁹ and approximately one-third of BC metastases are to the lungs.^{7,40} We previously showed that IL-2c and aPD-L1 fail as single-agent treatments in lung metastatic MB49 or MBT-2 BC, but their combination was effective and required CD8⁺ T cells.²⁰ To test if treatment mechanisms were cancer-

specific, we studied a model of melanoma lung metastasis via intravenous B16 cell injection. As in BC lung metastasis, aPD-L1 was unable to slow tumor growth (Figure 5a) or prolong mouse survival (Figure 5b) in lung metastatic B16 melanoma despite its efficacy in subcutaneous (Figure 3) and peritoneal (Fig. S7) metastases. However, IL-2c significantly slowed B16 lung metastatic tumor growth (Figure 5c) and improved survival as a single agent versus isotype treated controls (Figure 5d), mirroring outcomes in orthotopic (peritoneal) ID8agg ovarian cancer which is also IL-2c sensitive but aPD-L1 resistant.¹⁹

IL-2c treatment efficacy in lung metastasis is NK celldependent

Next, we tested if treatment mechanisms for IL-2c treatment efficacy in B16 lung metastases were similar to subcutaneous or peritoneal B16. Using TCR β^{KO} mice that lack CD4⁺ and CD8⁺ T cells, we found that IL-2c still reduced B16 lung tumor growth



Figure 5. IL-2c, but not α PD-L1, treats B16 lung metastases. Wild-type male mice were challenged intravenously with 3×10^5 B16 cells and treated with 100 µg α PD-L1 (a, b) on days 8, 13, 18, or IL-2c (c, d) on days 8, 10, 12, 14, or isotype control (a-c). (a,c) Tumor bioluminescence corresponding with pre-treatment, mid-treatment, and post treatment timepoints. N = 6–10 mice/group. p, two-way ANOVA of day 21 (a) or 20 (c) signal. (b,d) Mouse survival of mice in (a) and (c). N = 6–10 mice/group. p, log-rank. ANOVA, analysis of variance. IL-2c, interleukin-2 complex. PD-L1, Programmed Death Ligand 1.

(Figure 6a) and improved mouse survival versus isotype control treatment (Figure 6b). IL-2c acts through $\gamma\delta$ T cells in orthotopic bladder cancer.²⁰ To assess possible $\gamma\delta$ T cell effects in B16 lung metastases, we challenged RAG1^{KO} mice, which additionally lack $\gamma\delta$ T cells and B cells compared to TCR β^{KO} mice. IL-2c still improved mouse survival versus controls (Figure 6c), supporting the concept that IL-2c treats lung metastatic B16 independent of adaptive immunity or $\gamma\delta$ T cells.

As IL-2c significantly promoted NK cell activity in both orthotopic BC (Figure 1) and metastatic melanoma models (Figure 4), we tested NK cell-driven antitumor immunity as a mediator of IL-2c efficacy in metastatic B16 lung tumors. We depleted NK cells using α NK1.1 (Fig. S10) in mice bearing lung metastatic B16 melanoma treated with IL-2c, which reduced IL-2c-mediated control of tumor growth (Figure 6d) and survival (Figure 6e), supporting a critical role for NK cells in IL-2c efficacy here.



Figure 6. NK cells are essential for IL-2c efficacy in lung metastasis. $TCR\beta^{KO}$ (a, b), RAG1^{KO} (c), or wild-type (d,e) male mice were challenged intravenously with 3×10^5 B16. (a-e) Mice received IL-2c on days 8, 10, 12, 14, an isotype control (a-c), or 250 ug aNK1.1 on days 7, 10, 13, 16 (d, e). (a) Tumor bioluminescence in intravenous B16 challenge of $TCR\beta^{KO}$ mice. N = 9–10 mice/group. p, two-way ANOVA of day 20 signal. (b) $TCR\beta^{KO}$ mouse survival following intravenous B16 challenge. N = 9–10 mice/group. p, log-rank. (C) RAG1^{KO} mouse survival following intravenous B16 challenge. N = 9 mice/group. p, log-rank. (d) Tumor bioluminescence in intravenous B16 challenge of wild-type mice. N = 11–12 mice/group. p, two-way ANOVA of day 22 signal. (e) Wild-type mice survival following intravenous B16 challenge. Note: Survival of isotype-treated mice bearing B16 lung metastasis is in Figure 5d. N = 11–12 mice/group. p, log-rank. ANOVA, analysis of variance. RAG1, Recombinase Activating Gene 1. TCR, T Cell Receptor.

In lung metastatic MB49 and MBT-2 BC, IL-2c, or α PD-L1 alone are ineffective, but their combination effectively treats both models.²⁰ To test NK cell contributions to treatment of lung metastases in BC, we challenged mice with either lung metastatic MB49 or MBT-2 and treated with α PD-L1 + IL-2c with or without NK cell depletion (Fig. S11). In support of significant NK cell contributions, NK cell depletion significantly reduced mouse survival in both models (Fig. S12A,B). Together, these data show that IL-2c effects on NK cells are disparate from CD8⁺ T cell effects and contribute independently to immunotherapy efficacy across multiple primary (orthotopic) and metastatic models of BC and melanoma on distinct genetic backgrounds.

IL-2c activates lung NK cells to treat B16 lung metastases

We then performed flow cytometric analyses of B16 lung tumors (Fig. S13) to explore possible mechanisms for NK cell-mediated IL-2c efficacy in lung. We used gp100⁺ (melanoma-associated antigen⁴¹) cell number to normalize tumor-infiltrating NK and CD8⁺ T cell numbers and found that IL-2c but not aPD-L1, significantly increased the frequency (Figure 7a) and total numbers (Figure 7b) of lung tumor-infiltrating NK cells, as well as numbers of activated NK cells (Figure 7c). IL-2c also improved NK cell production of anti-tumor effector molecules granzyme B and IFN- γ (Figure 7d,e), and increased CD122 expression (figure 7f), with aPD-L1 having no observable NK cell effects (Figure 7a-f). Similar to orthotopic MB49 BC, IL-2c treatment also significantly increased expression of the NK cell maturation marker KLRG1 (Figure 7g), consistent with terminal differentiation and enhanced effector function of lung NK cells.

In contrast, neither IL-2c nor α PD-L1 treatment affected CD8⁺ T cell prevalence (Fig. S14A), total numbers (Fig. S14B), activation (Fig. S14 C), or effector molecule production (Fig. S14D). These results support our findings in T cell deficient mice (Figure 6a-c) that CD8⁺ T cells are dispensable for IL-2c efficacy in B16 lung metastasis and further emphasize the role of NK cells in mediating the IL-2c efficacy in this model.

Discussion

ICB with α PD-1 or α PD-L1 is FDA-approved to treat nearly 20 cancer subtypes⁴² yet even in the most ICB-sensitive cancers, many patients do not respond,⁴³ necessitating improved treatment approaches. Here, we studied IL-2c and α PD-L1 treatment mechanism differences in primary (orthotopic) and metastatic BC and melanoma, heading toward our goal of identifying conserved immune cell intrinsic mechanisms of these two agents that persist in varied tumor immune micro-environments, which could help define rational treatment combinations for improved treatment efficacy.

We previously reported that $\gamma\delta$ T cells were required for optimal IL-2c treatment efficacy in orthotopic bladder cancer²⁰ and recently showed that $\gamma\delta$ T cells augment antigen-specific immunity in the treatment of MB49 BC with Bacillus Calmette-Guérin (BCG) immune therapy.⁴⁴ Here, we found that NK cells also contribute to IL-2c efficacy in orthotopic bladder cancer in a $\gamma\delta$ T cell-independent manner, suggesting that both cells could contribute individually, rather than serially, to treatment efficacy. Thus, strategies to boost function of either cell type could improve IL-2c efficacy, such as ultralow dose rapamycin, which we demonstrated to be a safe and tolerable $\gamma\delta$ T cell adjuvant in early phase BC trials when combined with BCG immune therapy.⁴⁵

To understand tissue-selective effects and assess differences from aPD-L1, we found that aPD-L1 required both CD8⁺ T cells and NK cells to treat orthotopic B16 melanoma, whereas IL-2c only required CD8⁺ T cells.³⁴ These data show that IL-2c has differential mechanisms of action against the same tumor in distinct anatomic compartments, and that aPD-L1 effects in the same environment and tumor can differ from IL-2c. Our prior study showed that CD8⁺ T cells are dispensable for IL-2c treatment of orthotopic bladder cancer (where $\gamma\delta T$ cells²⁰ and NK cells (this report) are required), whereas aPD-L1 requires CD8⁺ T cells to treat orthotopic bladder cancer, suggesting potential benefit of combining both agents. We also found that NK cells from IL-2c treated mice consistently expressed lower PD-1, which could also impact ICB therapy. Further studies evaluating the impact of CD122-directed IL-2 therapy on ICB targets (e.g., PD-1, PD-L1) are warranted to gain better mechanistic insights that can facilitate optimized combination therapies.

We considered that metastatic tumors represent an additional immune challenge to hosts, whereby the tumor appears in distinct anatomic compartments. To understand IL-2c and aPD-L1 effects in metastatic tumors, we evaluated peritoneal B16 melanoma, a large unmet clinical need, as patients with peritoneal metastatic melanoma have a median survival of only two months.⁸ We found that either IL-2c or aPD-L1 alone effectively treated, but did not cure, peritoneal B16. By contrast, we previously showed that IL-2c very effectively treated peritoneal (orthotopic) ID8agg ovarian cancer, where aPD-L1 is entirely ineffective.¹⁹ Thus, these outcomes do not reflect the generic efficacy of aPD-L1 in the peritoneal compartment. In peritoneal B16, IL-2c activated CD8⁺ T cells more effectively versus NK cells, which could be due to greater CD122 expression on local CD8⁺ T cells versus NK cells that we show here. Cell depletion data were consistent with flow data for aPD-L1 efficacy mechanisms, but not for IL-2c. IL-2c greatly improved NK cell numbers and functions over aPD-L1, yet NK cell depletion very surprisingly did not affect IL-2c treatment efficacy in peritoneal B16. There are several potential explanations that require further investigations. Among these, the NK cell effects required for treatment efficacy might not be reflected in flow analyses, specific suppressive factors could impede NK cell efficacy after IL-2c but not after aPD-L1, and NK cell/tumor co-colocalization could differ based on IL-2c versus aPD-L1 treatments. These data underscore the fact that flow cytometric analyses in this model might not fully reflect all mechanistic details. Further, as CD8⁺ T cell depletion did not fully reverse IL-2c treatment efficacy in peritoneal metastases, additional mechanisms might exist, requiring further studies.

In the clinically relevant setting of B16 melanoma lung metastases, IL-2c outperformed α PD-L1 in tumor control and host survival. As α PD-L1 treats subcutaneous B16, these data show that the local immune microenvironment can significantly affect treatment efficacy. In accord with much other data here, IL-2c efficacy in lung metastases depended on NK cells. Further, lung metastatic IL-2c efficacy was independent of



Figure 7. IL-2c promotes CD122 expression and NK cell maturation in B16 lung metastases. Wild-type male mice were challenged intravenously with 3×10^5 B16 cells and treated with 100 µg αPD-L1 on days 14, 19, or IL-2c on days 14, 16, 18, or isotype control, and sacrificed on day 20. Total lung NK cell frequency (a) and tumornormalized NK cell number (b), CD69 expression (c), granzyme B production (d), IFN- γ production (e), CD122 expression (f), and KLRG1 expression (g). N = 5–9 mice/ group. *p* value, one-way ANOVA. ANOVA, analysis of variance. IFN- γ , interferon-gamma. IL-2c, interleukin-2 complex. KLRG1, Killer Cell Lectin Like Receptor G1. MFI, Mean Fluorescence Intensity. NK, Natural Killer. PD-L1, Programmed Death Ligand 1.

CD8⁺ T cells (or tumor-specific immunity), as we found that IL-2c efficacy was preserved in RAG^{KO} mice lacking any tumor-specific immunity. Flow cytometry analyses of B16 lung tumors showed that IL-2c preferentially boosts lung NK cell activation and effector function but not lung CD8⁺ T cells,

confirming the necessity of NK cells in IL-2c treatment of B16 lung metastasis and providing a striking contrast to peritoneal B16 tumors where IL-2c activated both CD8⁺ T and NK cells. We also identified an increase in lung KLRG1⁺ NK cells following IL-2c treatment, which denotes a terminally mature NK cell population with a protective effect against pulmonary metastasis.^{46,47} The robust effects on this population in both bladder and lung following IL-2c treatment warrants further investigation, as IL-2c-driven NK cell maturation may occur via a CD122-Eomes axis,⁴⁸ with expansion of KLRG1⁺ NK cells serving as a possible biomarker for responsiveness to CD122-directed immunotherapies. In this regard, our NK cell immunophenotyping generally agrees with prior reports.²⁹ However, as tissue-specific NK cell differences are known,⁴⁹ detailed NK cell studies in specific anatomic compartments with tumor metastases will help further understanding of immunotherapy effects. Thus, detailed understandings of the immune land-scape in distinct anatomic compartments and effects of specific treatment agents therein will help define more effective treatment regimens.

All together, we establish differential responsiveness of the same tumor to the same agent in distinct (metastatic) compartments, and IL-2c mechanisms of efficacy appear largely to depend on NK cells, whereas aPD-L1 effects depend to a greater extent on CD8⁺ T cells. These data suggest that combining aPD-L1 with IL-2c could be more effective against a variety of cancers, including in metastatic settings where there is a great unmet medical need. Further, other CD122-selective treatment approaches under evaluation, such as pegylated IL-2 (bempegaldesleukin) which shows efficacy in metastatic melanoma trials in combination with the aPD-1 agent, nivolumab,⁵⁰ should be similarly tested. It remains to be seen whether distinct CD122-targeted IL-2 constructs have differing capacities to activate $CD8^+$ T cells versus NK cells or $\gamma\delta$ T cells, but such data will greatly help define optimal combinations for further clinical evaluations.

Abbreviations

Analysis of variance (ANOVA), asialo GM1 (asGM1), Bacillus Calmette-Guérin (BCG), bladder cancer (BC), cytotoxic T-lymphocyte-associate protein 4 (CTLA-4), Eomes (Eomesodermin), IL-2 complex (IL-2c), IL-2 receptor (IL-2R), immune checkpoint blockade (ICB), interleukin (IL), interferon-gamma (IFN- γ), killer cell lectin-like receptor G1 (KLRG1), mean fluorescence intensity (MFI), milligram (mg), natural killer (NK), programmed death receptor 1 (PD-1), programmed death-ligand 1 (PD-L1), recombination activating gene 1 (RAG1), T regulatory cells (Tregs), T cell receptor (TCR)

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Consent for publication

All authors of this manuscript consent to its submission and publication.

Availability of data and material

The datasets used and/or analyzed during the current study are available from the corresponding author upon reasonable request.

Authors' contributions (CRediT – Contributor Roles Taxonomy)

RMR: conceptualization, data curation, formal analysis, funding acquisition, investigation, methodology, project administration, validation, visualization, writing – original draft, writing – review and editing CZ: data curation, formal analysis, investigation, methodology, visualization, writing – original draft, and writing – review and editing. YD: data curation, formal analysis, investigation, methodology, visualization, and writing – review and editing. NJ: investigation, methodology, resources. NM: data curation, methodology, resources. ASP: investigation, formal analysis, funding acquisition, resources, supervision, and writing – review and editing. TJC: conceptualization, data curation, formal analysis, funding acquisition, methodology, project administration, resources, software, supervision, writing – original draft, writing – review and editing.

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