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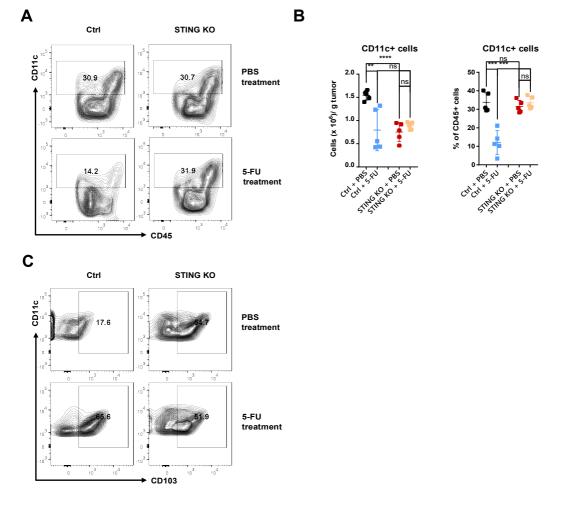
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sgRNA sequences		
STING sgRNA 1	mouse	GTAGAATCATAGCCATACAG
STING sgRNA 2	mouse	GAAGGCCAAACATCCAACTG
STING sgRNA 3	mouse	CAGTAGTCCAAGTTCGTGCG
cGAS sgRNA 1	mouse	CAGAATGCAGAAACGGGAGT
cGAS sgRNA 2	mouse	GAGGCGCGGAAAGTCGTAAG
Ifnb1 sgRNA 1	mouse	GAAACAATTTCTCCAGCACT
Ifnb1 sgRNA 2	mouse	GTTTCTGAAGACAGTACTAG
Ifna1 sgRNA 1	mouse	TCACATCCTAGAGAGCAGGT
Ifna1 sgRNA 2	mouse	GAAGATGTTCAGGATCTGCT
Ifnar1 sgRNA 1	mouse	TTCAGCAGAATATCGAACGT
Ifnar1 sgRNA 2	mouse	CTTCTAAACGTACTTCTGGG

Appendix Table S1. sgRNA sequences used in this study.

RT primers		
lfnb1-f	mouse	CCCTATGGAGATGACGGAGA
lfnb1-r	mouse	CCCAGTGCTGGAGAAATTGT
Irf7-f	mouse	AGAGGCGTTTTATCTTGCG
lfr7-r	mouse	TGGAGCCCAGCATTTTCTCT
Stat1-f	mouse	CACGCTGCCTATGATGTCTC
Stat1-r	mouse	ACGCTTGCTTTTCCGTATGT
Ifit1-f	mouse	CAGAAGCACACATTGAAGAA
lfit1-r	mouse	TGTAAGTAGCCAGAGGAAGG
Gapdh-f	mouse	ATGTGTCCGTCGTGGATCTGAC
Gapdh-r	mouse	AGACAACCTGGTCCTCAGTGTAG

Appendix Table S2. Primers used in this study.

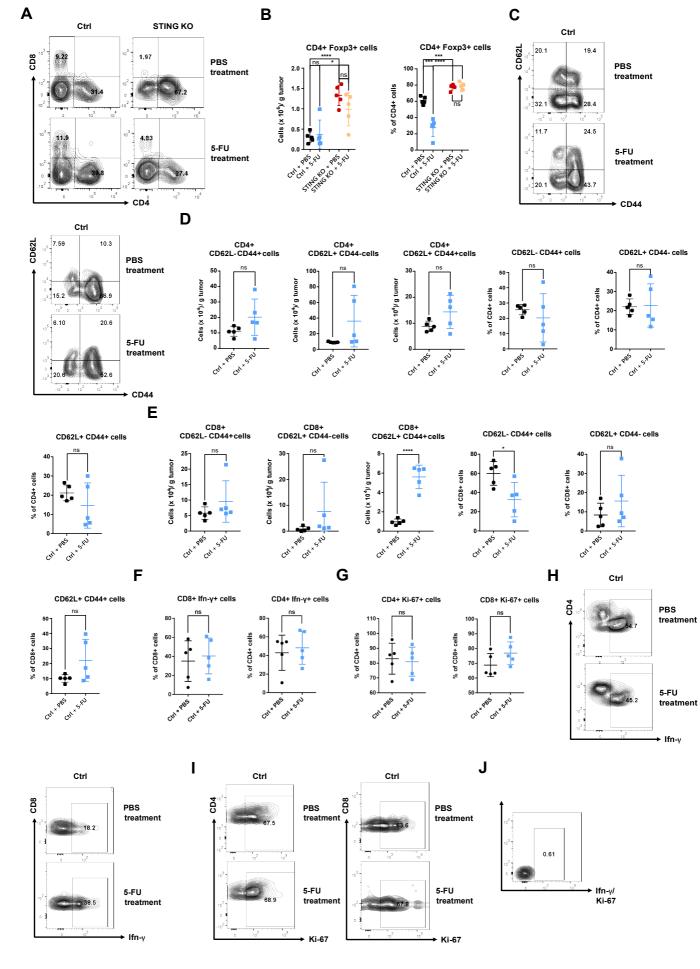


Tian et al. Appendix Figure S1

Appendix Figure S1. Effect of 5-FU treatment on intratumoral dendritic cells.

- **A.** Representative flow cytometry plots showing the percentages of intratumoral CD11c⁺ cells, after gating on CD45⁺ cells.
- **B.** The counts of intratumoral CD11c⁺ cells per gram of tumor were quantified (left panel). The percentages of CD11c⁺ cells within intratumoral CD45⁺ cells (right panel). Each dot represents one mouse. N=5.
- **C.** Representative flow cytometry plots showing the percentages of CD103⁺ cells within intratumoral CD11c⁺ cells.

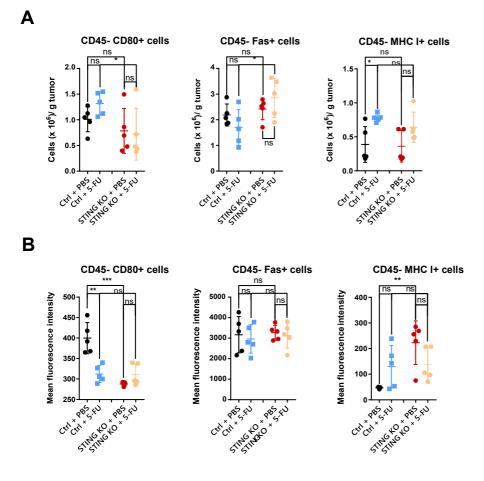
For all panels, error bars represent s.d., and center values represent mean. Two-tailed unpaired student's t-test was used. **: p < 0.01; ***: p < 0.001; ***: p < 0.0001; ns: not significant.



Tian et al. Appendix Figure S2

Appendix Figure S2. Analysis of intratumoral T cells after 5-FU treatment

- **A.** Representative flow cytometry plots showing the percentages of CD4⁺ or CD8⁺ T cells within intratumoral CD3⁺ cells.
- **B.** (Left) The counts of intratumoral Tregs (CD4⁺FoxP3⁺) per gram of tumor were quantified. (Right) The percentages of Tregs among CD4⁺ cells were quantified. N=5. Each dot represents one mouse.
- **C.** Representative flow cytometry plots analyzing intratumoral CD4+ (left panel) or CD8+ (right panel) T cells with CD62L and CD44.
- **D.** The counts of CD4⁺ intratumoral CD62L⁻ CD44⁺, CD62L⁺ CD44⁻, and CD62L⁺ CD44⁺ cells per gram of tumor were quantified. Additionally, the corresponding percentage of each population among CD4⁺ T cells were quantified. N=5. Each dot represents one mouse.
- **E.** The counts of CD8⁺ intratumoral CD62L⁻ CD44⁺, CD62L⁺ CD44⁻, and CD62L⁺ CD44⁺ cells per gram of tumor were quantified. Additionally, the corresponding percentage of each population among CD8⁺ T cells were quantified. N=5. Each dot represents one mouse.
- **F-J.** Control MC38 cells were injected into mice and mice were treated with 5-FU or PBS. Tumors were harvested at 2 weeks post cancer cell injection. **(F)** The percentages of intratumoral Ifn- γ ⁺ cells within CD4⁺ (left panel) or CD8⁺ (right panel) cells were quantified. **(G)** The percentages of intratumoral Ki67⁺ cells within CD4⁺ (left panel) or CD8⁺ (right panel) cells were quantified. Each dot represents one mouse. **(H)** Representative flow cytometry plots showing the percentages of Ifn- γ ⁺ cells within CD4⁺ and CD8⁺ cells. **(I)** Representative flow cytometry plots showing the percentages of Ki67⁺ cells within CD4⁺ and CD8⁺ cells. **(J)** Flow cytometry plots showing the negative control of staining for (H) and (I). For all panels, error bars represent s.d., and center values represent mean. Two-tailed unpaired student's t-test was used. *: p < 0.05; ****: p < 0.0001; ns: not significant.



Tian et al. Appendix Figure S3

Appendix Figure S3. Effect of 5-FU on surface antigens in intratumoral CD45⁻ cells.

A-B. Mice were injected with control (Ctrl) or STING-KO MC38 cells and treated with PBS or 5- FU. Tumors were harvested 2 weeks after cancer cell injection and intratumoral CD45⁻ cells were examined by flow cytometry. **(A)** The counts of CD45⁻CD80⁺, CD45⁻Fas⁺, and CD45⁻MHC-I⁺ cells per gram of tumor were quantified. Each dot represents one mouse. N=5. **(B)** The mean fluorescence intensities of CD80, Fas and MHC-I in CD45⁻CD80⁺, CD45⁻Fas⁺, and CD45⁻MHC-I⁺ cells, respectively, were quantified. Each dot represents one mouse. N=5.

For all panels, error bars stand for s.d., and center values represent mean. Two-tailed unpaired student's t-test was used. *: p < 0.05; ns: not significant. Data are representative of two independent experiments.