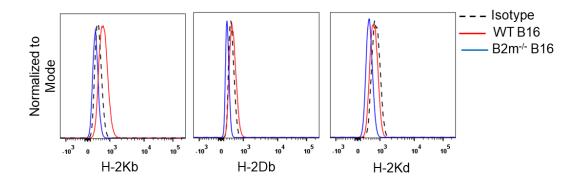
# **Supplementary Material**

Title: Natural killer cells suppress cancer metastasis by eliminating circulating cancer cells

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## **SUPPLEMENTARY FIGURES**



**Figure S1. MHC-I expression on WT and B2m**<sup>-/-</sup> **B16 cells.** The expression of MHC-I alloantigens H-2Kb (C57BL/6), H-2Db (C57BL/6) and H-2Kd (BALB/c) were examined on the surface of WT and B2m<sup>-/-</sup> B16 melanoma cells.

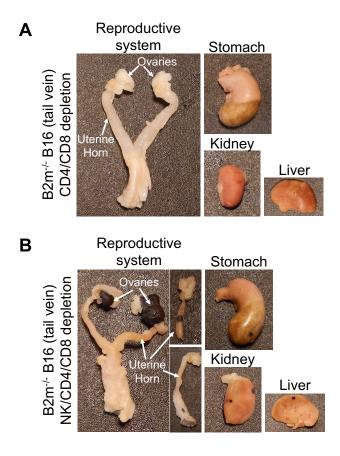


Figure S2. NK and T cell depletion results in extrapulmonary metastases of B16 melanoma.

(A, B) Representative macroscopic images of extrapulmonary metastases of B2m<sup>-/-</sup> B16 melanoma following tail vein IV injection and treatment with anti-CD4/CD8 (A) and anti-NK1.1/CD4/CD8 (B) antibodies.

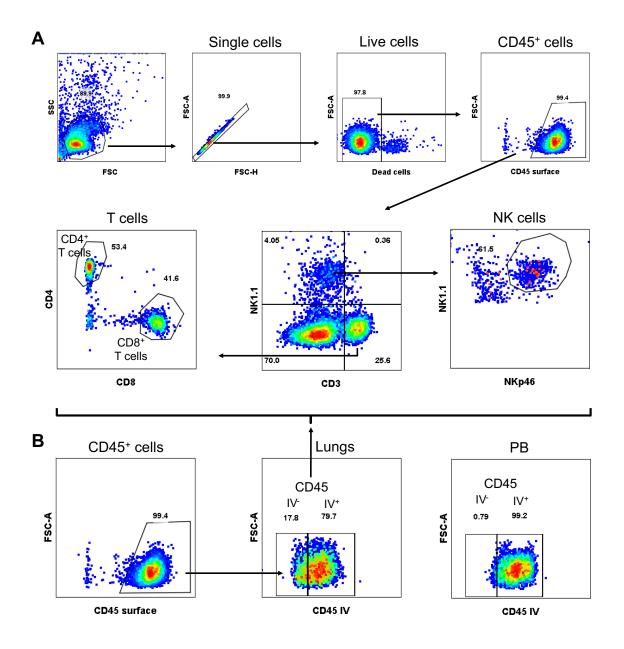


Figure S3. Gating strategy to detect total and tissue infiltrating (IV<sup>neg</sup>) NK and T cells. For specific evaluation of the immune cells infiltrating into the lung tissue, fluorescently labeled anti-CD45 antibody was intravenously administered to mice 3 minutes before organ harvest. CD45-stained immune cells from the vasculature are termed as CD45 IV<sup>+</sup> while tissue-resident/emigrated immune cells which remained unstained by intravenously injected CD45 antibody are termed as CD45 IV<sup>-</sup>. Following the organ harvest, immune cells were *ex vivo* stained by distinctly labeled CD45 antibody ("CD45 surface") to stain all immune cells. (A) Total CD45<sup>+</sup>

immune cells gated among forward/side scatter-based singlets gate followed by fixable live/dead stain negative live cells gate. Total NK cells were subsequently detected by gating for CD3<sup>-</sup>, NK1.1<sup>+</sup>, NKp46<sup>+</sup> population. Total CD4<sup>+</sup> and CD8<sup>+</sup> T cells were detected by gating for NK1.1<sup>-</sup>, CD3<sup>+</sup>, CD4<sup>+</sup> or CD8<sup>+</sup> populations. (B) To detect circulating versus tissue-infiltrating immune cells in the lung, total *ex vivo* labeled CD45<sup>+</sup> immune cells were gated for CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) populations accounting for circulating and tissue-infiltrating immune cells. The gating of CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) immune cells among total CD45<sup>+</sup> cells was generated based on the peripheral blood (PB) sample where all CD45<sup>+</sup> immune cells were CD45 IV<sup>+</sup> (IV<sup>pos</sup>) (i.e., fluorescently labeled by intravenously injected CD45 antibody). CD45 IV<sup>-</sup> (IV<sup>neg</sup>) NK cells and CD4<sup>+</sup>/CD8<sup>+</sup> T cells were further detected as described in (A).

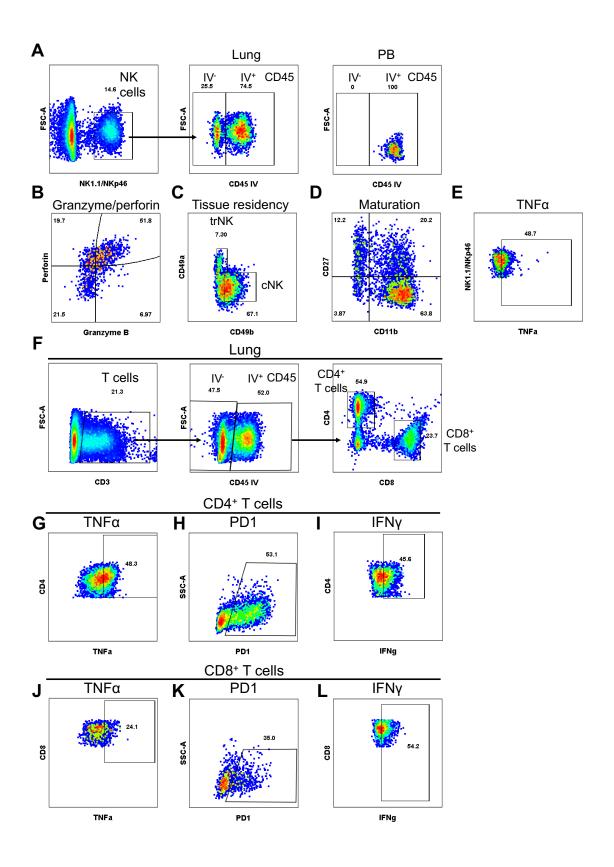


Figure S4. Gating strategy for functional characterization of Lung IV<sup>pos</sup> and IV<sup>neg</sup> NK and T cells. (A) NK cells detected by gating for CD3<sup>-</sup>, NK1.1/ NKp46<sup>+</sup> population. To detect circulating versus tissue-infiltrating immune cells, mice were intravenously injected with fluorescently labeled CD45 antibody 3 minutes before organ harvest to fluorescently label all CD45<sup>+</sup> immune cells in the vasculature. The gating of CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) NK cells among total NK cells was generated based on the peripheral blood (PB) sample where all NK cells were CD45 IV<sup>+</sup> (IV<sup>pos</sup>) (i.e., fluorescently labeled by intravenously injected CD45 antibody). (B-E) CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) lung NK cells gated for granzyme B/perforin (B), CD49b/CD49a (C), CD11b/CD27 (D), and TNFα (E) based on the Fluorescence Minus One (FMO) controls. (F) T cells detected by gating for CD3<sup>+</sup>, NK1.1/ NKp46<sup>-</sup> population. The gating of CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) T cells was generated as for NK cells and were subsequently gated for CD4<sup>+</sup> and CD8<sup>+</sup> T cells. (G-L) CD45 IV<sup>+</sup> (IV<sup>pos</sup>) and CD45 IV<sup>-</sup> (IV<sup>neg</sup>) lung CD4<sup>+</sup> (G, H, I) and CD8<sup>+</sup> (J, K, L) T cells were further gated for TNFα (G, J), PD1 (H, K), and IFNγ (I, L) based on the FMO controls.

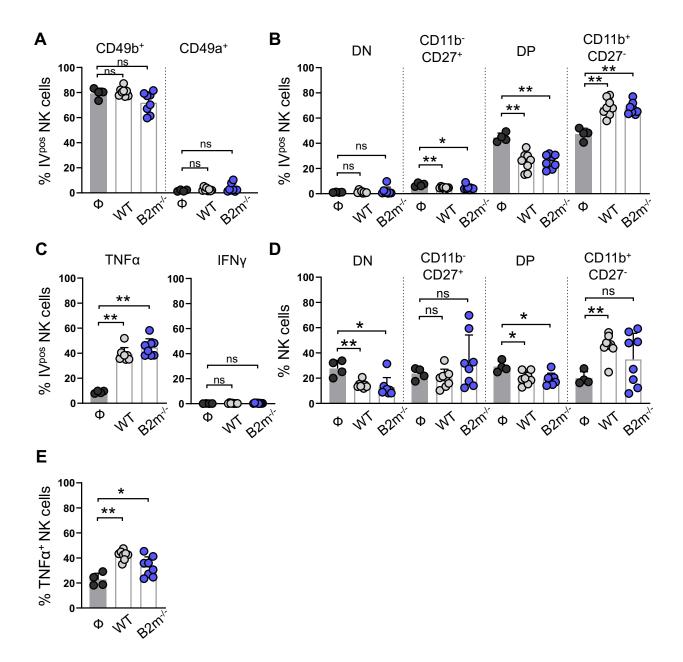


Figure S5. Characterization of lung and liver NK cells in response to metastatic melanoma.

(A) Quantification of conventional (CD49b<sup>+</sup>) and tissue-resident (CD49a<sup>+</sup>) circulating (IV<sup>pos</sup>) NK cells in the lung at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice (Φ). (B) Quantification of maturation status defined by CD11b and CD27 expression in circulating (IV<sup>pos</sup>) NK cells in the lung at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice

(Φ). **(C)** Quantification of TNF $\alpha^+$  and IFN $\gamma^+$  circulating (IV<sup>pos</sup>) NK cells in the lung at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice (Φ). **(D)** Quantification of maturation status defined by CD11b and CD27 expression in liver NK cells at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice (Φ). **(E)** Quantification of TNF $\alpha^+$  liver NK cells at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice (Φ). n = 4-8 mice per group. Graphs show mean + SD. Mann-Whitney *U* test, ns: not significant, \* p < 0.05, \*\* p < 0.01.

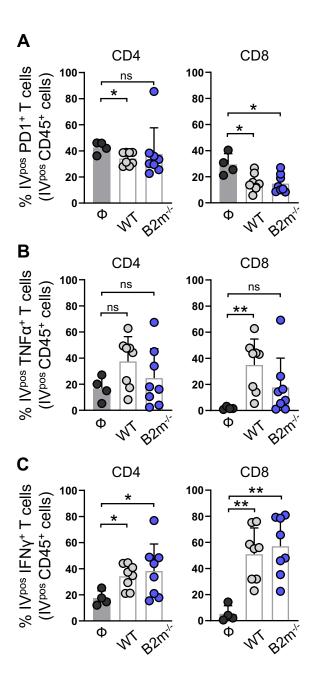


Figure S6. Characterization of circulating T cells in the lung in response to metastatic melanoma. (A-C) Quantification of PD1<sup>+</sup> (A), TNFα<sup>+</sup> (B), and IFNγ<sup>+</sup> (C) circulating (IV<sup>pos</sup>) CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the lung at day 14 after WT B16 (WT) and B2m<sup>-/-</sup> B16 (B2m<sup>-/-</sup>) melanoma tail vein IV injection in comparison with untreated mice (Φ). n = 4-8 mice per group. Graphs show mean + SD. Mann-Whitney *U* test, ns: not significant, \* p < 0.05, \*\* p < 0.01.

#### **SUPPLEMENTARY TABLE**

Table S1. Antibodies used for immune cell analysis.

| <u>Clone</u> | <u>Company</u>                         |
|--------------|--|
| CD3-12       | Abcam, Cambridge, MA                   |
| D4W2Z        | Cell Signaling Technology, Danvers, MA |
| EPR19514     | Abcam                                  |
|              | CD3-12                                 |

## **Secondary Antibodies and staining kits**

Goat anti-Rat IgG, Alexa Fluor® 568 conjugate

Goat anti-Rabbit IgG, Alexa Fluor® 488

conjugate

Thermo Fisher Scientific, Waltham, MA

Thermo Fisher Scientific

# **Flow Cytometry Antibodies**

| Mouse   |  |   |
|---|--|---|
| CD3-BUV395  | 145-2C11   | BioLegend   |
| CD3-APC-Cy7   | 145-2C11   | BioLegend   |
| CD3-PerCP-Cy5.5   | 145-2C11   | BioLegend   |
| CD4-APC-Cy7   | GK1.5  | BioLegend   |
| CD4-eF450   | GK1.5  | BioLegend   |
| CD4-PerCP-Cy5.5   | GK1.5  | BioLegend   |
| CD8-PE  | 53-6.7   | BioLegend   |
| CD8-APC-Cy7   | 53-6.7   | BioLegend   |
| CD8-BUV395  | 53-6.7   | BioLegend   |
| CD45-BV605  | 30-F11   | BioLegend   |
| CD45-APC  | 30-F11   | BioLegend   |
| CD27-PE-Cy7   | LG.3A10  | BioLegend   |
|   | LG.SATU  | DioLogona   |
| Perforin-PE   | S16009A  | BioLegend   |
| •   |  | •   |
| Perforin-PE   | S16009A  | BioLegend   |
| Perforin-PE<br>CD19-APC-Cy7   | S16009A<br>6D5   | BioLegend<br>BioLegend  |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5  | S16009A<br>6D5<br>Ha31/8   | BioLegend<br>BioLegend<br>BioLegend   |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5 CD49b-FITC   | S16009A<br>6D5<br>Ha31/8<br>DX5                                      | BioLegend<br>BioLegend<br>BioLegend<br>BioLegend                                |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5 CD49b-FITC NK1.1-APC                                     | S16009A<br>6D5<br>Ha31/8<br>DX5<br>PK136                             | BioLegend BioLegend BioLegend BioLegend BioLegend                               |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5 CD49b-FITC NK1.1-APC NK1.1-PECy7                         | S16009A<br>6D5<br>Ha31/8<br>DX5<br>PK136<br>PK136                    | BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend                     |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5 CD49b-FITC NK1.1-APC NK1.1-PECy7 NK1.1-APC-Cy7           | S16009A<br>6D5<br>Ha31/8<br>DX5<br>PK136<br>PK136<br>PK136           | BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend           |
| Perforin-PE CD19-APC-Cy7 CD49a-PerCP-Cy5.5 CD49b-FITC NK1.1-APC NK1.1-PECy7 NK1.1-APC-Cy7 NK946-APC | S16009A<br>6D5<br>Ha31/8<br>DX5<br>PK136<br>PK136<br>PK136<br>29A1.4 | BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend BioLegend |

| PD1-FITC                | 29F.1A12 | BioLegend   |
|-------------------------|----------|-------------|
| IFNγ-eF450              | XMG1.2   | eBioscience |
| IFNγ-AF700              | XMG1.2   | BioLegend   |
| IFNγ-PE-Cy7             | XMG1.2   | eBioscience |
| IFNγ-BV605              | XMG1.2   | BioLegend   |
| TNFα-BV605              | MP6-XT22 | BioLegend   |
| Granzyme B-AF647        | GB11     | BioLegend   |
| Granzyme B-Pacific Blue | GB11     | BioLegend   |
| H-2Kb-APC               | AF6-88.5 | BioLegend   |
| H-2Db-PE                | KH95     | BioLegend   |
| H-2Kd-PE-Cy7            | SF1-1.1  | BioLegend   |