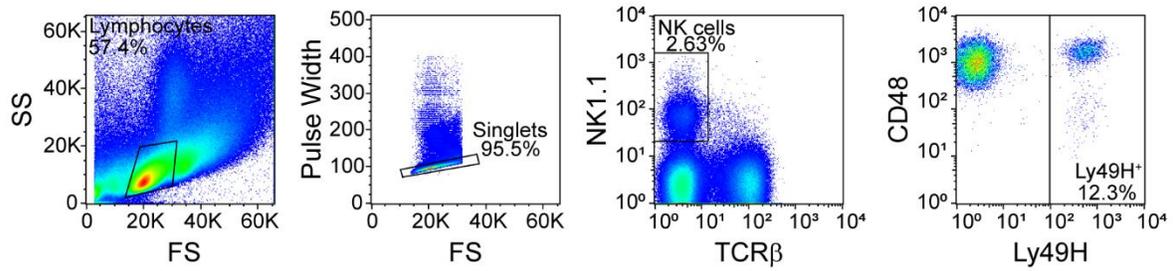
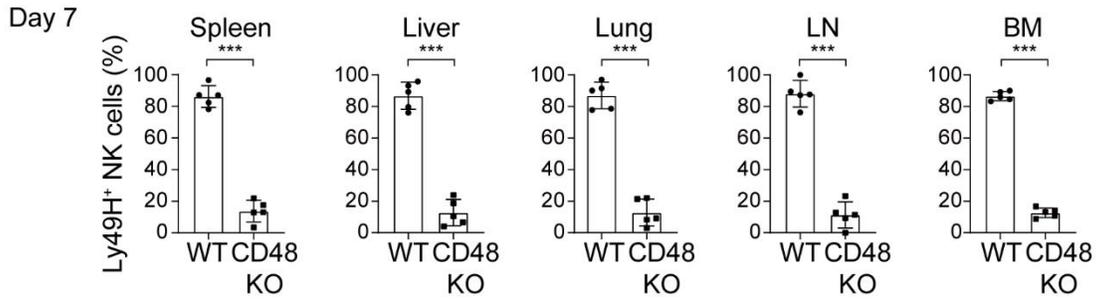
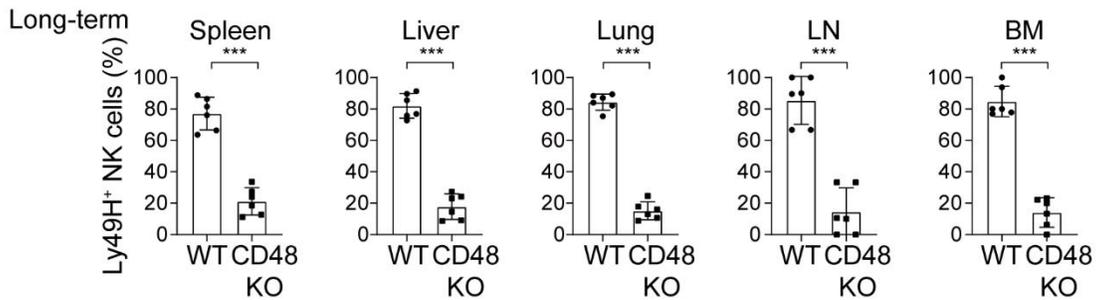
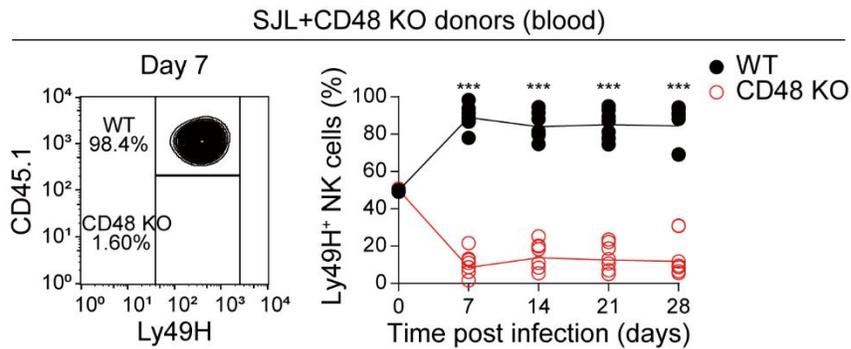
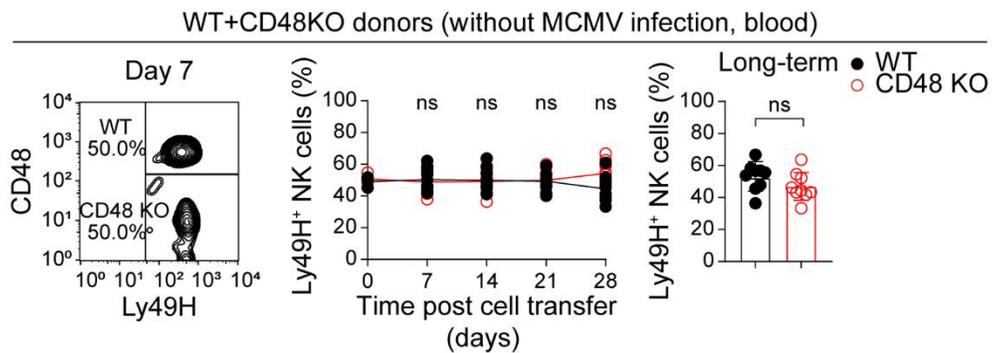


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**Supplemental information**

**Suppression of adaptive NK cell expansion  
by macrophage-mediated phagocytosis  
inhibited by 2B4-CD48**

**Rui Li, Cristian Camilo Galindo, Dominique Davidson, Huaijian Guo, Ming-Chao Zhong, Jin Qian, Bin Li, Zsolt Ruzsics, Colleen M. Lau, Timothy E. O'Sullivan, Silvia M. Vidal, Joseph C. Sun, and André Veillette**

**A****B****C****D****E**

**Figure S1. CD48 deficiency selectively impairs MCMV-induced expansion of Ly49H<sup>+</sup> NK cells. Related to Figure 1 and Figure 2.**

(A) Representative flow cytometry analysis of the general gating strategy of Ly49H<sup>+</sup> NK cells (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>, Ly49H<sup>+</sup>) used in this study is depicted.

(B) Same as Figure 2B, except the proportions of WT and CD48 KO Ly49H<sup>+</sup> NK cells were measured in various organs, including the spleen, liver, lung, lymph node (LN) and bone marrow (BM) at day 7 (n = 5).

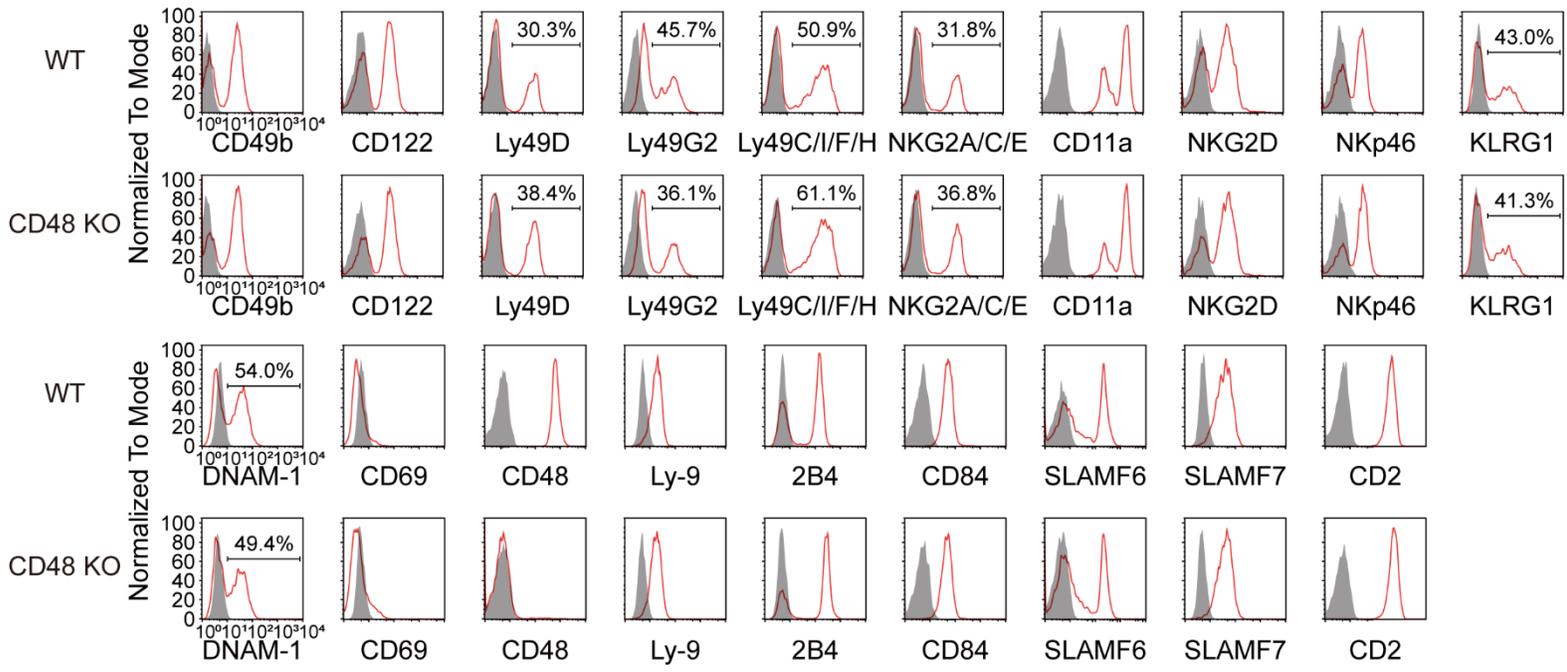
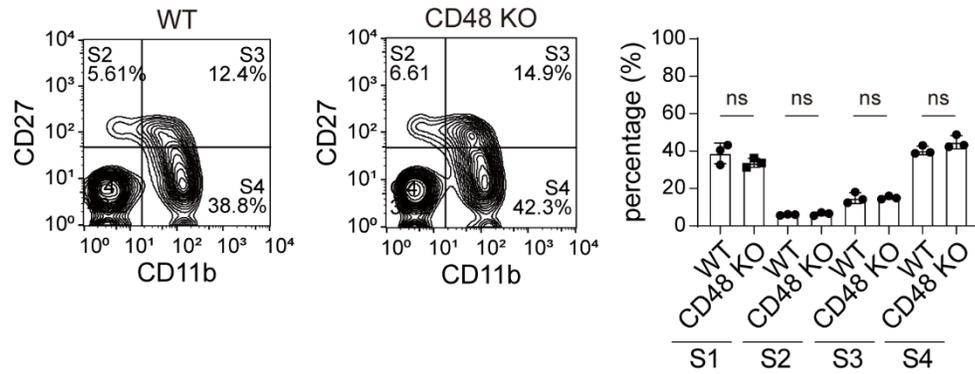
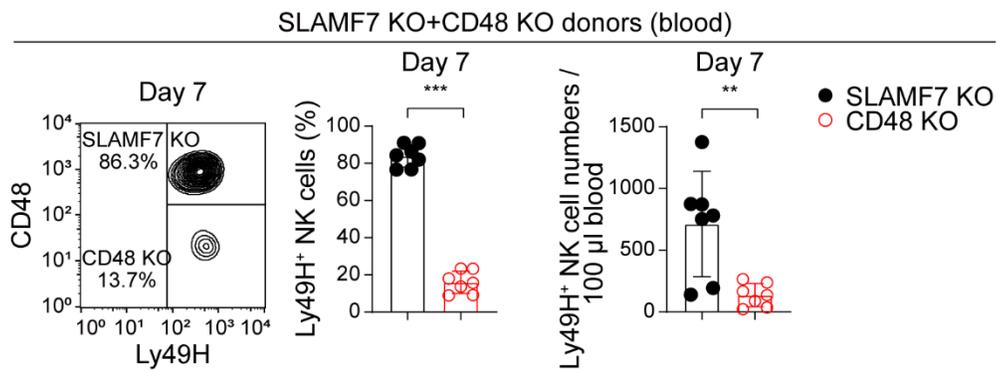
(C) Same as (B), except the results on day 56 are depicted (n = 6).

(D) Blood. Equal numbers of Ly49H<sup>+</sup> NK cells from WT(B6.SJL; CD45.1<sup>+</sup>/CD45.2<sup>-</sup>) and CD48 KO (CD45.1<sup>-</sup>/CD45.2<sup>+</sup>) donors were mixed and injected into Ly49H KO mice. The proportions of WT versus CD48 KO NK cells in the Ly49H<sup>+</sup> population (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>, Ly49H<sup>+</sup>) in blood of recipient mice were analyzed at indicated time points. CD45.1 was used as a marker to distinguish WT (CD45.1<sup>+</sup>) and CD48 KO (CD45.1<sup>-</sup>) cells. Left panel: representative flow cytometry analysis of CD45.1 expression on Ly49H<sup>+</sup> NK cells at day 7. Proportions of WT (CD45.1<sup>+</sup>) and CD48 KO (CD45.1<sup>-</sup>) Ly49H<sup>+</sup> NK cells are depicted. Right panel: proportions of WT and CD48 KO Ly49H<sup>+</sup> NK cells at various time points in multiple independent mice (n = 7).

(E) Same as Figure 2B, except the adoptive transfer experiments were performed without MCMV infection. From left to right: first panel, representative flow cytometry analysis of CD48 expression on Ly49H<sup>+</sup> NK cells in blood of recipient mice at day 7. Proportions of

WT (CD48<sup>+</sup>) and CD48 KO (CD48<sup>-</sup>) Ly49H<sup>+</sup> NK cells are depicted. Second panel: proportions of WT and CD48 KO Ly49H<sup>+</sup> NK cells at various time points in multiple independent mice (n = 9). Third panel: proportions of WT and CD48 KO Ly49H<sup>+</sup> NK cells at day 35 in blood for multiple independent mice (n = 9).

Statistical analyses were conducted using unpaired t-tests (two-tailed), except in panels displaying data with multiple time points, where multiple t-tests were employed. Each symbol represents an individual mouse; error bars depict the mean with s.d. \*p ≤ 0.05, \*\*p ≤ 0.01, \*\*\*p ≤ 0.001, ns (not significant). Data are representative of 2 (B); 2 (C); 2 (D) and 3 (E) independent experiments, two to four recipient mice were injected with purified NK cells in each independent experiment.

**A****B****C**

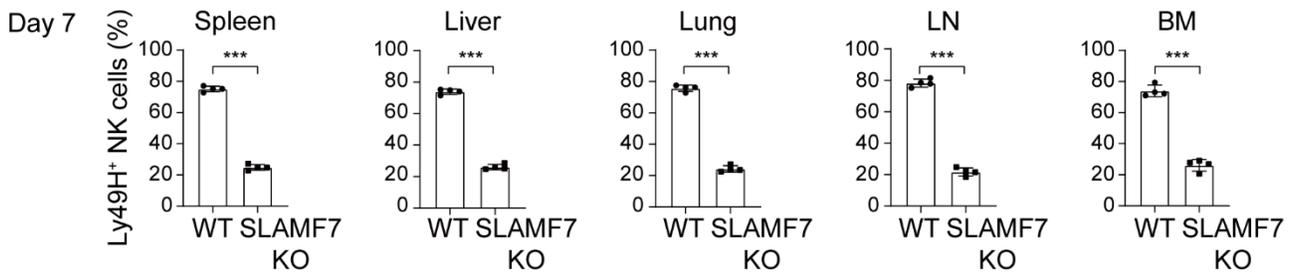
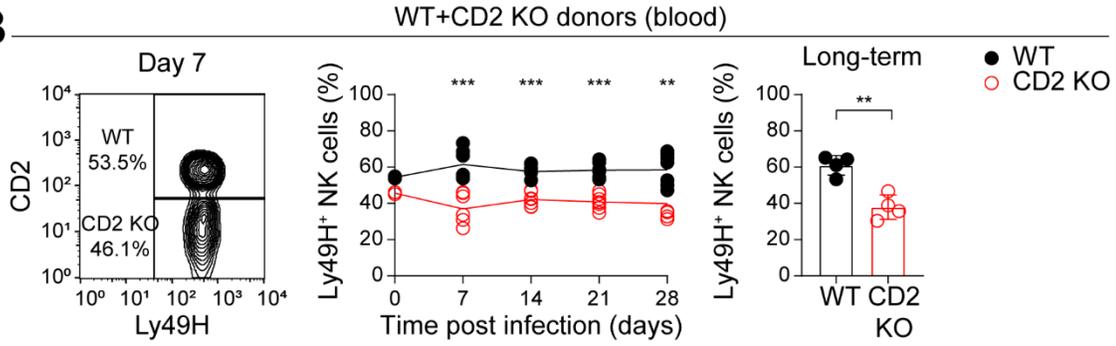
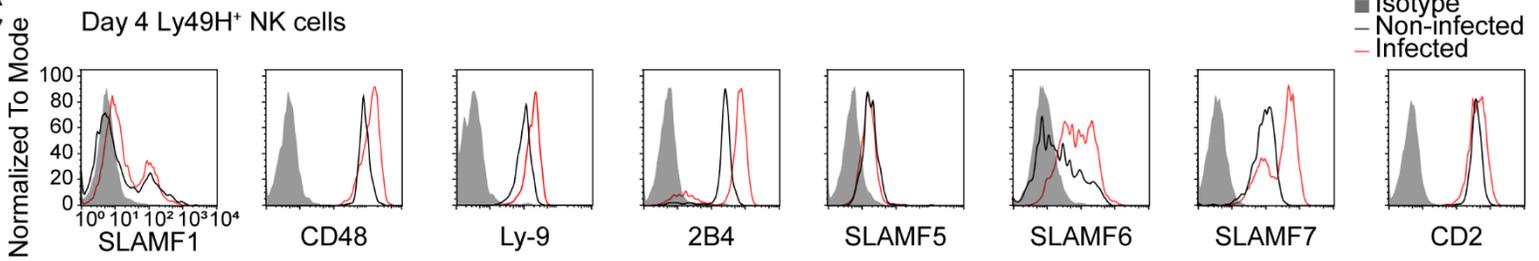
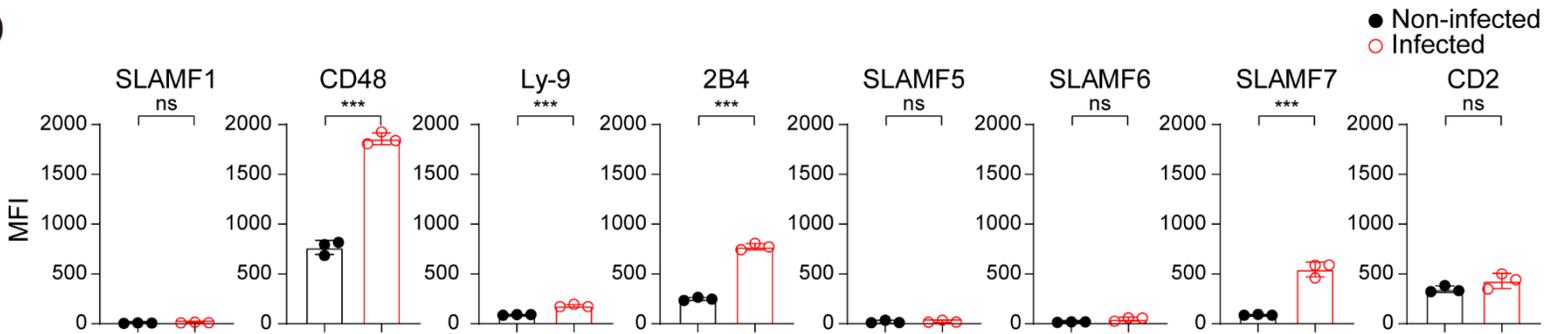
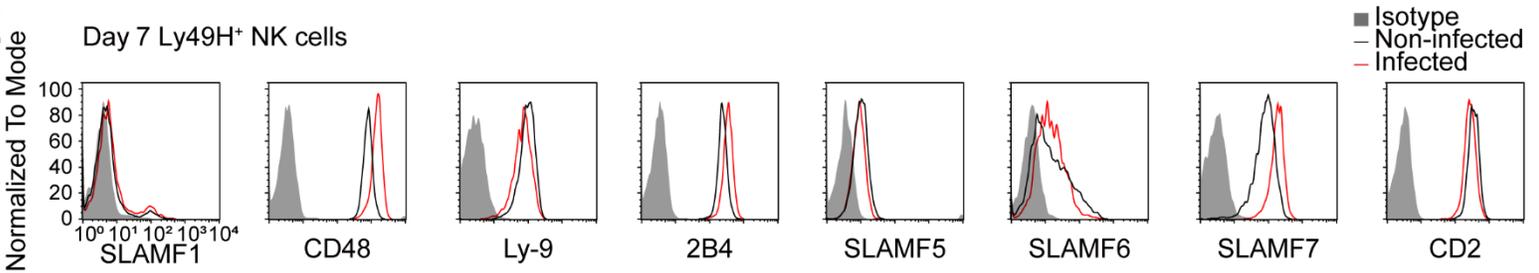
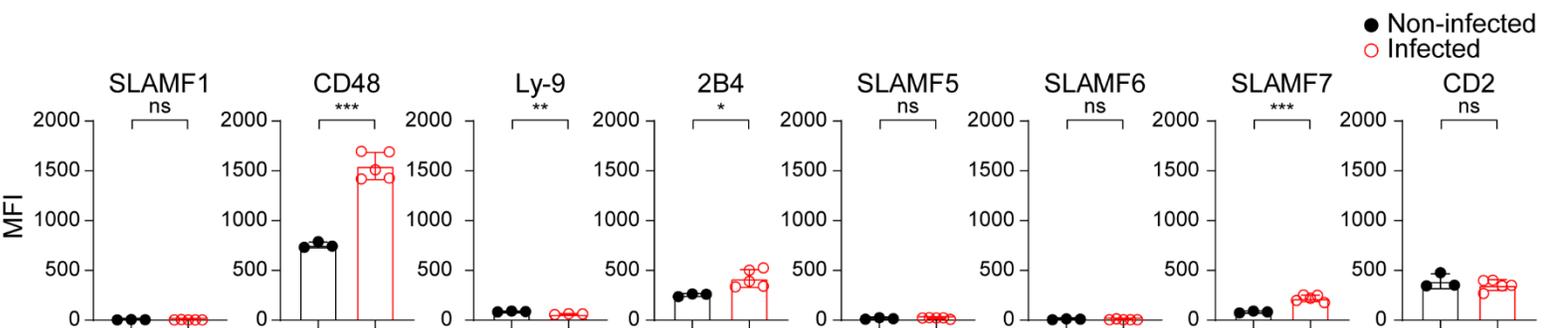
**Figure S2. CD48 deficiency does not affect NK cell development. Related to Figure 2.**

(A) Representative flow cytometry analyses of NK cell receptor repertoires on freshly isolated splenic NK cells (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>) from WT or CD48 KO mice (n = 3).

(B) Flow cytometry analyses of development of freshly isolated splenic NK cells (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>) from WT or CD48 KO mice. CD11b and CD27 were stained to determine the development status of NK cells. Stage 1 (S1), CD11b<sup>-</sup>/CD27<sup>-</sup>; Stage 2 (S2), CD11b<sup>-</sup>/CD27<sup>+</sup>; Stage 3 (S3), CD11b<sup>+</sup>/CD27<sup>+</sup>; Stage 4 (S4), CD11b<sup>+</sup>/CD27<sup>-</sup>. From left to right: first two panels: representative analyses of CD11b and CD27 expression on WT and CD48 KO NK cells. Third panel: proportions of various developmental stages for WT and CD48 KO NK cells (n = 3).

(C) Equal numbers of Ly49H<sup>+</sup> NK cells from SLAMF7 KO and CD48 KO donors were mixed and injected into Ly49H KO mice. The proportions of SLAMF7 KO versus CD48 KO NK cells in the Ly49H<sup>+</sup> population (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>, Ly49H<sup>+</sup>) in blood of recipient mice were analyzed at day 7. CD48 was used as a marker to distinguish SLAMF7 KO (CD48<sup>+</sup>) and CD48 KO (CD48<sup>-</sup>) cells. From left to right: first panel, representative flow cytometry analysis of CD48 expression on Ly49H<sup>+</sup> NK cells at day 7. Proportions of SLAMF7 KO (CD48<sup>+</sup>) and CD48 KO (CD48<sup>-</sup>) Ly49H<sup>+</sup> NK cells are depicted. Second panel: statistical analysis of the proportions for multiple independent mice (n = 7). Third panel: numbers of SLAMF7 KO and CD48 KO Ly49H<sup>+</sup> NK cells per 100 μl blood (n = 7).

Statistical analyses were conducted using unpaired t-tests (two-tailed). Each symbol represents an individual mouse; error bars depict the mean with s.d. \* $p \leq 0.05$ , \*\* $p \leq 0.01$ , \*\*\* $p \leq 0.001$ , ns (not significant). Data are representative of 3 independent experiments.

**A****B****C****D****E****F**

**Figure S3. CD48 plays a more crucial role than SLAMF7 in the expansion of Ly49H<sup>+</sup> NK cells. Related to Figure 2.**

(A) Same as Figure 2C, except the proportions of WT and SLAMF7 KO Ly49H<sup>+</sup> NK cells were measured in various organs including the spleen, liver, lung, lymph node (LN) and bone marrow (BM) at day 7 (n = 4).

(B) Blood. Equal numbers of Ly49H<sup>+</sup> NK cells from WT and CD2 KO donors were mixed and injected into Ly49H KO mice. The proportions of WT versus CD2 KO NK cells in the Ly49H<sup>+</sup> population (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>, Ly49H<sup>+</sup>) in blood of recipient mice were analyzed at indicated time points. CD2 was used as a marker to distinguish WT (CD2<sup>+</sup>) and CD2 KO (CD2<sup>-</sup>) cells. From left to right: first panel, representative flow cytometry analysis of CD2 expression on Ly49H<sup>+</sup> NK cells at day 7. Proportions of WT (CD2<sup>+</sup>) and CD2 KO (CD2<sup>-</sup>) Ly49H<sup>+</sup> NK cells are depicted. Second panel: proportions of WT and CD2 KO Ly49H<sup>+</sup> NK cells at various time points in multiple independent mice (n = 6). Third panel: proportions of WT and CD2 KO Ly49H<sup>+</sup> NK cells in blood at day 56 (n = 4).

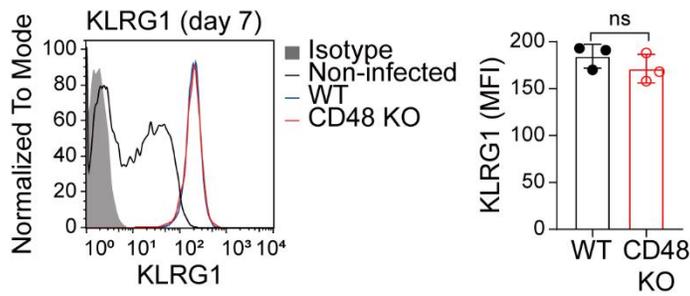
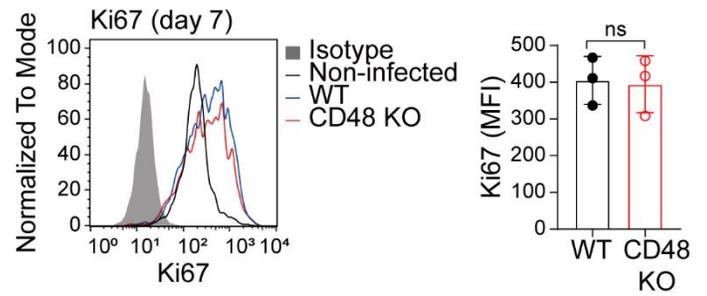
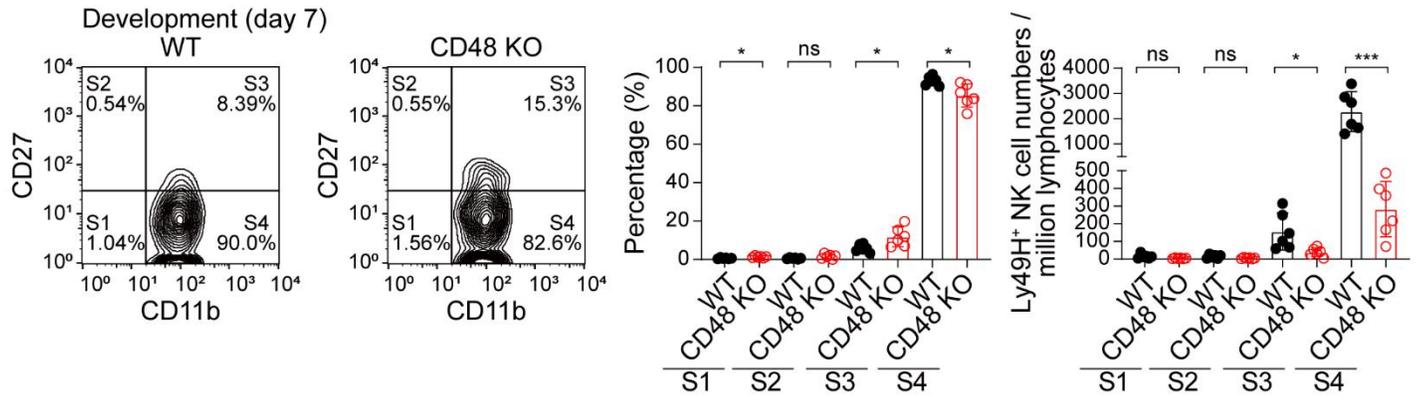
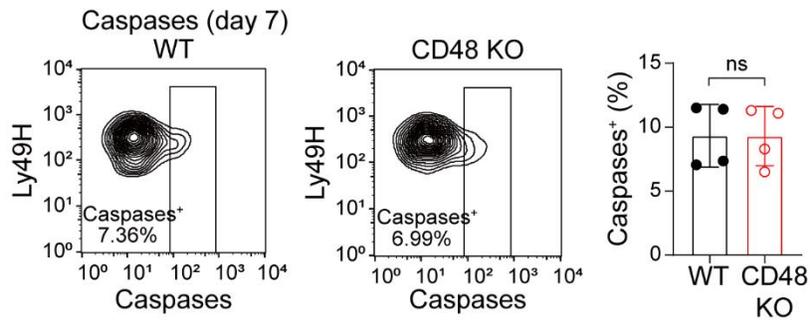
(C) On day -1 (1 day before infection), Ly49H<sup>+</sup> NK cells from WT donors were injected into Ly49H KO mice. On day 0, 7.5 x 10<sup>2</sup> pfu of MCMV were injected intra-peritoneally into recipients. The SLAM family receptors and CD2 expression on Ly49H<sup>+</sup> NK cells (NK1.1<sup>+</sup>, TCRβ<sup>-</sup>, Ly49H<sup>+</sup>) in spleen of recipient mice were analyzed at day 4. Isotype controls are shown in grey, non-infected controls are shown in black, and the cells from

infected mice are shown in red. Representative flow cytometry analyses of the expression of the receptors on Ly49H<sup>+</sup> donor NK cells at day 4 are depicted.

(D) Same as (C), except the statistical analyses of flow cytometry results for multiple mice are depicted (n = 3).

(E and F) Same as (C and D), except flow cytometry analysis results at day 7 are depicted. n = 3-5.

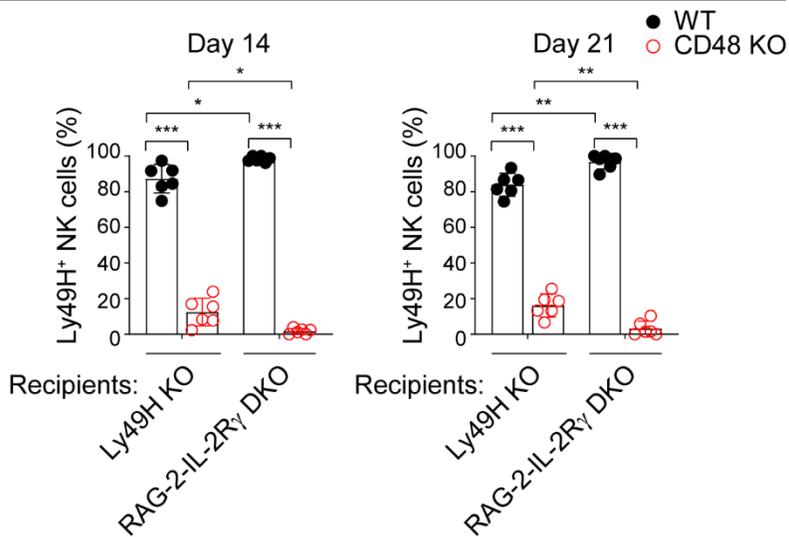
Statistical analyses were conducted using unpaired t-tests (two-tailed), except in panels displaying data with multiple time points, where multiple t-tests were employed. Each symbol represents an individual mouse; error bars depict the mean with s.d. \*p ≤ 0.05, \*\*p ≤ 0.01, \*\*\*p ≤ 0.001, ns (not significant). Data are representative of 2 (A); 2 (B) and 3 (C-F) independent experiments, one to four recipient mice were injected with purified NK cells in each independent experiment..

**A****B****C****D**

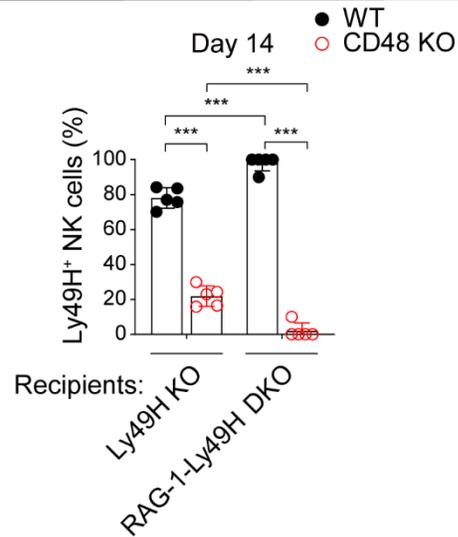
**Figure S4. Minimal effect of CD48 deficiency on NK cell activation and proliferation.**

**Related to Figure 3.**

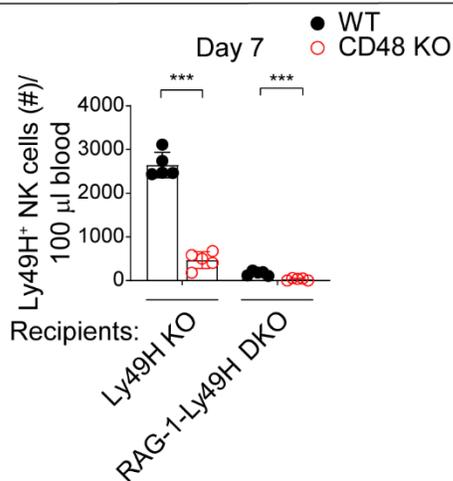
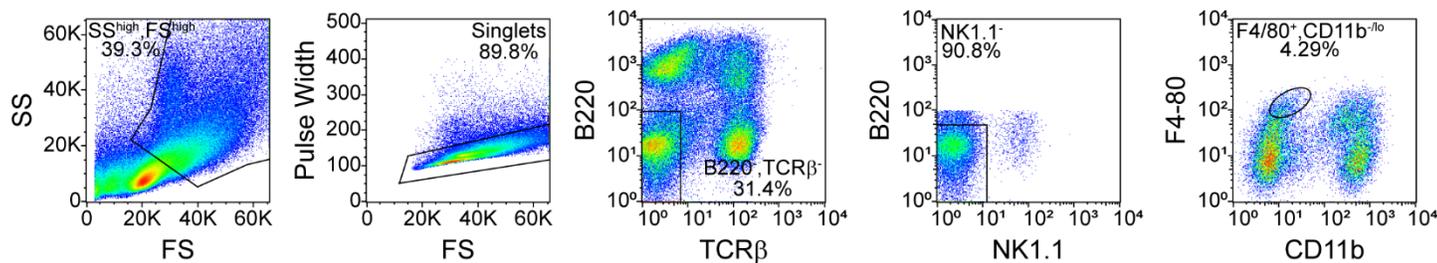
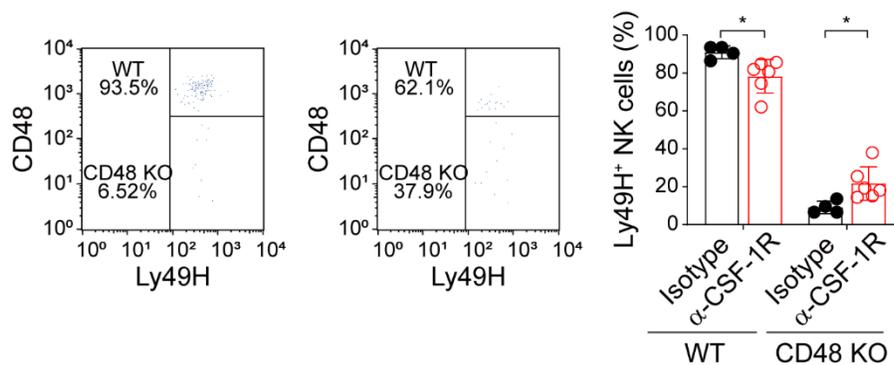
(A-D) Same as in Figure 3B-E, except results at day 7 are depicted. Statistical analyses were conducted using unpaired t-tests (two-tailed). Each symbol represents an individual mouse; error bars depict the mean with s.d. \* $p \leq 0.05$ , \*\* $p \leq 0.01$ , \*\*\* $p \leq 0.001$ , ns (not significant).

**A**WT+CD48 KO donors; RAG-2-IL-2R $\gamma$  DKO recipients (blood)**B**

WT+CD48 KO donors; RAG-1-Ly49H DKO recipients (blood)

**C**

WT+CD48 KO donors; RAG-1-Ly49H DKO recipients (blood)

**D****E**Macrophage depletion ( $\alpha$ -CSF-1R, blood)

**Figure S5. Macrophages limit the expansion of Ly49H<sup>+</sup> NK cells. Related to Figure 5.**

(A) Same as in Figure 5B, except results at day 14 and day 21 are depicted (n = 6).

(B) Same as in Figure 5C, except results at day 14 are depicted (n = 5).

(C) Same as in Figure 5C, except numbers of WT and CD48 KO Ly49H<sup>+</sup> NK cells per 100  $\mu$ l blood at day 7 are depicted (n = 5).

(D) Representative flow cytometry analysis of the general gating strategy of fresh isolated macrophages (B220<sup>-</sup>, TCR $\beta$ <sup>-</sup>, NK1.1<sup>-</sup>, F4/80<sup>+</sup>, CD11b<sup>-/lo</sup>) used in this study is depicted.

(E) Same as Figure 5E, except blood was analyzed [n = 4 (isotype control); n = 6 ( $\alpha$ -CSF-1R)].

Statistical analyses were conducted using unpaired t-tests (two-tailed), except in panels that involved comparisons of more than two groups, where one-way ANOVA followed by Tukey's multiple comparison tests were employed to assess significance. Each symbol represents an individual mouse; error bars depict the mean with s.d. \*p  $\leq$  0.05, \*\*p  $\leq$  0.01, \*\*\*p  $\leq$  0.001, ns (not significant). Data are representative of 2 independent experiments, two to three recipient mice were injected with purified NK cells in each independent experiment.