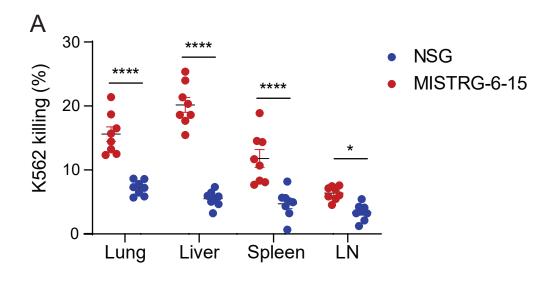
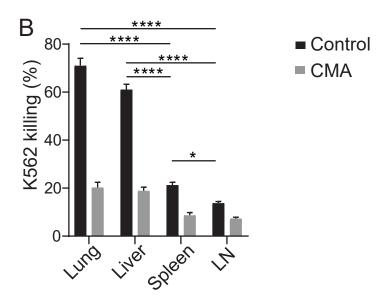


Supplemental Fig. 1. Immune cell distribution in MISTRG-6-15 mice.

A, Representative flow of CD14 and NKp46 staining in spleen, lung and liver. **B,** Percentage of human T cells (CD3), NK cells (NKp46), and macrophages (CD14) in spleen, lung, and liver. Results from several independent cohorts were combined.

Figure S2

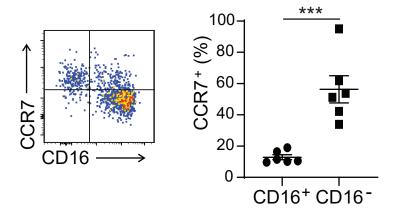




Supplemental Fig. 2. K562 killing by NK cells from NSG and MISTRG-6-15 mice.

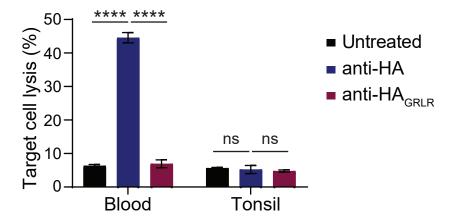
A, ex vivo killing of K562 by purified tissue NK cells after 4 hours of co-culture. **B**, ex vivo killing of K562 by purified tissue NK cells from MISTRG-6-15 mice with and without concanamycin A. NK cells and K562 cells were co-cultured at 1:1 for 4 hours before flow cytometry analysis. Error bars show mean values with SEM. P-values were calculated using a two-way ANOVA with Tukey multi-comparison post-test. * denotes p-value <0.05, ** denotes p-value <0.01, *** denotes p-value <0.001, and **** denotes p-value <0.0001.

Figure S3



Supplemental Fig. 3. CCR7 and CD16 expression in NK cells.

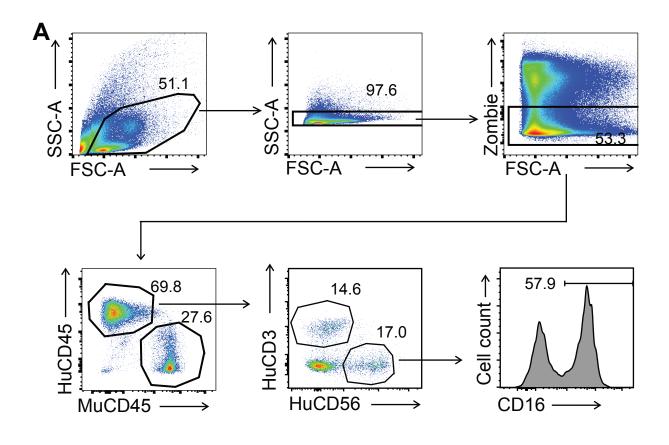
Representative flow plot of CCR7 and CD16 expression in blood NK cells. Percentage of CCR7 $^+$ in CD16 $^+$ and CD16 $^-$ NK cells. Error bars show mean values with SEM. Unpaired, two-tailed t-test used. *** denotes p-value <0.001.

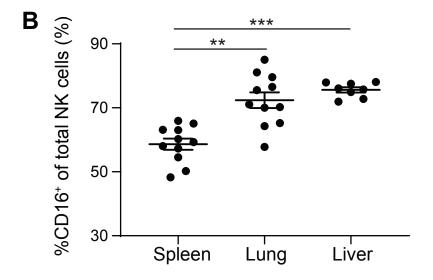


Supplemental Fig. 4. Killing of HIV-1-infected CD4⁺ cells by NK cells from blood and tonsil tissues.

CD4⁺ T cells infected with HIVivo-HA virus were co-cultured with autologous NK cells from either blood or tonsils as noted at 1:1 for 4 hours with the presence of anti-HA or anti-HA_{GRLR} (1μg/ml). Target cells lysis was determined by the percentage of lysed infected cells (CD3 +CD8-HA⁺7-AAD⁺) in the total infected cells (CD3+CD8-HA⁺). Error bars show mean values with SEM. *P*-values were calculated using a two-way ANOVA with Tukey multi-comparison post-test. **** denotes p-value <0.0001.

Figure S5

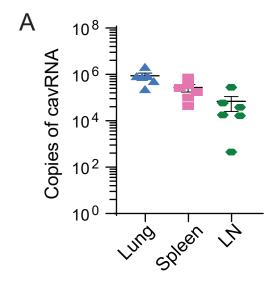


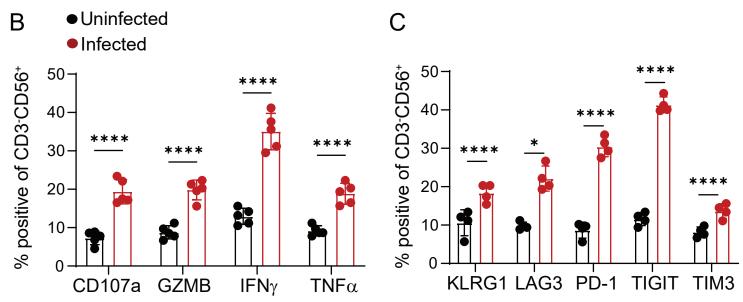


Supplemental Fig. 5. CD16⁺ NK cells in MISTRG-6-15 mice.

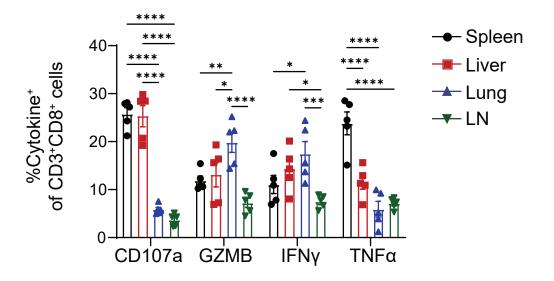
A, Human NK cell gating strategy. Total splenocytes were used. **B,** percentage of CD16⁺ NK cells in spleen, lung and liver. Error bars show mean values with SEM. *P*-values were calculated using a one-way ANOVA with Tukey multi-comparison post-test. ** denotes p-value <0.01 and *** denotes p-value <0.001.

Figure S6





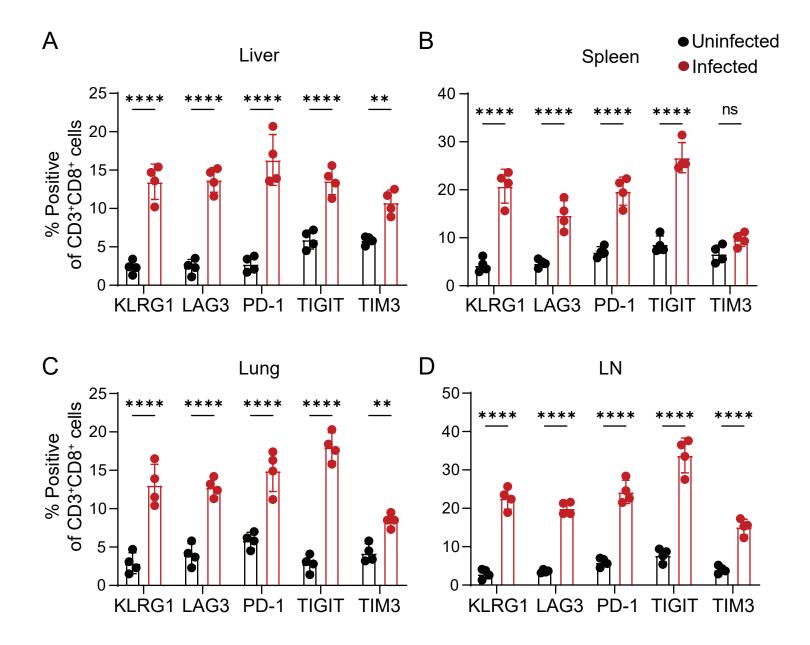
Supplemental Fig. 6. HIV-1 infection and blood NK responses. MISTRG-6-15 mice were infected with HIV-1_{BAL}. Blood and tissues were collected on day 21 post infection. A, Copies of cell-associated viral RNA (cavRNA) in lung, spleen and LN. B, Percentage of blood NK cells positive for CD107a, GZMB, IFNγ, and TNFα after ex vivo stimulation with PMA/ionomycin for 4 hours. C, Percentage of blood NK cells positive for KLRG1, LAG3, PD-1, TIGIT, and TIM3. Error bars show mean values with SEM. P-values were calculated using two-way ANOVA with Sidak's multiple comparison test. * denotes p-value <0.05, and **** denotes p-value <0.0001.



Supplemental Fig. 7. Functionality of CD8⁺ T cells during HIV-1 infection in different organs.

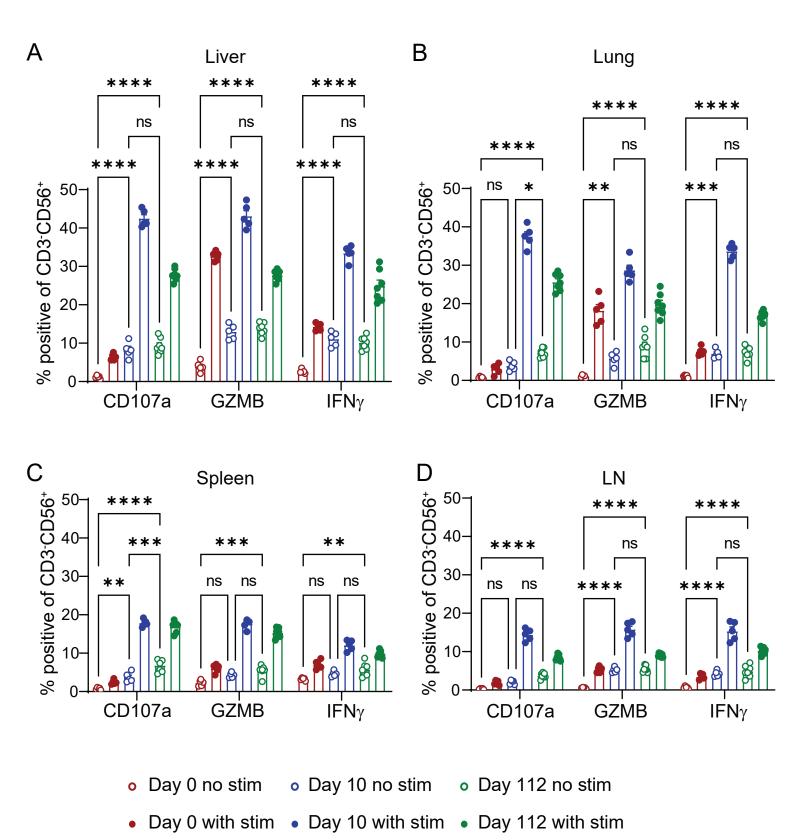
MISTRG-6-15 mice were infected with HIV-1_{BAL}. Tissues were collected on day 21 post infection. Percentage of CD8⁺ T cells positive for either CD107a, granzyme B, IFN γ , or TNF α in spleen, liver, lung, and lymph nodes after ex vivo stimulation with PMA/ionomycin for 4 hours. CD8⁺ T cells were gated on huCD45⁺CD3⁺CD8⁺. Error bars show mean values with SEM. *P*-values were calculated using two-way ANOVA with Tukey multi-comparison post-test. * denotes p-value <0.05, ** denotes p-value <0.01, *** denotes p-value <0.001, and **** denotes p-value <0.0001.

Figure S8

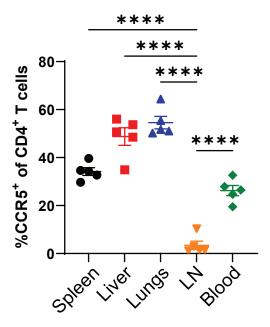


Supplemental Fig. 8. ICR expression in CD8⁺ T cells during HIV-1 infection in different organs.

MISTRG-6-15 mice were infected with HIV-1_{BAL}. Tissues were collected on day 21 post infection. Percentage of tissue CD8⁺ T cells positive for KLRG1, LAG3, PD-1, TIGIT, and TIM3 in liver, spleen, lung, and lymph nodes. CD8⁺ T cells were gated on huCD45⁺CD3⁺CD8⁺. Error bars show mean values with SEM. *P*-values were calculated using two-way ANOVA with Sidak's multiple comparison test. ** denotes p-value <0.01, and **** denotes p-value <0.0001.

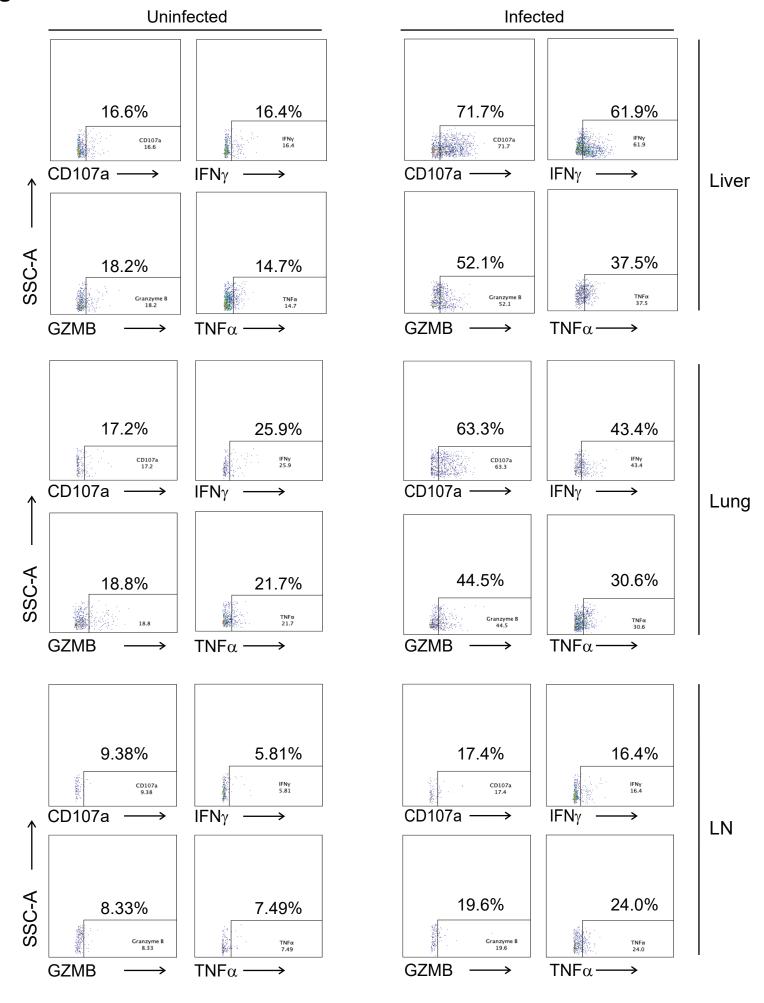


Supplemental Fig. 9. NK cell response throughout the course of HIV-1 infection. MISTRG-6-15 mice were infected with HIV-1_{BAL}. Tissue analyses were performed on day 0, 10, and 112 post infection. 4-6 mice at each time point were included. Percentage of CD107a⁺, GZMB⁺, and IFNγ⁺ NK cells in liver, lung, spleen, and LN with or without ex vivo stimulation with PMA/ionomycin for 4 hours. NK cells were gated on huCD45⁺CD3⁻CD56⁺. Error bars show mean values with SEM. *P*-values were calculated using two-way ANOVA with Tukey multicomparison post-test. ** denotes p-value <0.001, *** denotes p-value <0.0001.



Supplemental Fig. 10. CCR5⁺CD4⁺ T cells in tissues.

Percentage of CCR5⁺ of total CD4⁺ T cells in spleen, liver, lung, LN, and blood of MISTRG-6-15 mice. MISTRG-6-15 mice were infected with HIV-1_{BAL}. Tissue analyses were performed on day 14 post infection. Error bars show mean values with SEM. *P*-values were calculated using one-way ANOVA with Tukey multi-comparison post-test. **** denotes p-value <0.0001.



Supplemental Fig. 11. NK cell cytokine production by intracellular staining. MISTRG-6-15 mice were infected with HIV-1_{BAL} or left uninfected. Tissue samples were collected on day 21 post infection. Representative flow plots for CD107a, GZMB, IFN γ , and TNF α staining after ex vivo stimulation with PMA/ionomycin for 4 hrs. NK cells were gated on huCD45⁺CD3⁻CD56⁺.