Cell Reports, Volume 39

Supplemental information

Nasally delivered interferon- λ protects

mice against infection by SARS-CoV-2

variants including Omicron

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Figure S1. SARS-CoV-2 viral burden in infected K18-hACE2 mice, Related to Figure 2. (A) Eight-week-old female K18-hACE2 mice were inoculated by intranasal route with 10^3 FFU of WA1/2020 D614G. At -16 h before virus inoculation, mice were given 2 µg of murine IFN- λ 2 or PBS by intraperitoneal injection. Viral RNA levels at 3 dpi (n = 6-7 per group, 2 experiments). (**B-**G) Eight-week-old female K18-hACE2 mice were inoculated by intranasal route with 10^3 FFU of WA1/2020 D614G. At -16 h (**B-C**), D-3 (**D-E**) or +8 h (**F-G**), mice were given 2 µg of murine IFN- λ 2 or PBS by intranasal route. Viral RNA (**B, D, and F**) and infectious virus (**C, E, and G**) levels at 3 dpi (**B-C**: n = 7 per group, 2 experiments; **D-E**: n = 8-9 per group, 2 experiments; **F-G**: n = 6-7 per group, 2 experiments). (**H-J**) Eight-week-old female K18-hACE2 mice were treated with 2 µg doses of murine IFN- λ 2 or PBS by intranasal route IFN- λ 2 or PBS by intranasal route if N- λ 2 or PBS by intranasal route if N- λ 2 or PBS by intranasal route. (**H-J**) Eight-week-old female K18-hACE2 mice were treated with 2 µg doses of murine IFN- λ 2 or PBS by intranasal route at -16 h and +8 h relative to inoculation with 10^3 FFU of WA1/2020 D614G and harvested at 7 dpi. (**H**) Weight change was

monitored daily for 7 days. (I) Viral RNA levels at 7 dpi. (J) Infectious virus levels at 7 dpi (H-J: n = 9-10 per group, 2 experiments). Bars (A-G and I-J) indicate median values. Data were analyzed by Mann-Whitney test (A-G and I-J) or *t* tests of the area under the curve (H) (**P* < 0.05, ***P* < 0.01, ****P* < 0.001, and *****P* < 0.0001).



Figure S2. SARS-CoV-2 viral burden in the brains of K18-hACE2 and 129S2 mice, Related to Figures 2 and 3. (A-D) Eight-week-old (A-B) or five-month-old (C-D) female K18-hACE2 mice were inoculated by intranasal route with 10^3 FFU of WA1/2020 D614G (A-B) or B.1.1529 Omicron variant (C-D). At D-2 (A), D+1 and D+2 (B and D) or D-1 (C), mice were administered 2 µg of murine IFN- λ 2 or PBS by intranasal route. Viral RNA levels from brain at 3 dpi (A: n = 9 per group, 2 experiments; B: n = 8 per group, 2 experiments; C: n = 7-8 per group, 2 experiments; D: n = 6-7 per group, 2 experiments). (E-H) Six-week-old female 129S2 mice were inoculated by intranasal route with 10^5 FFU of B.1.351 Beta variant. At D-1 (E), D-3 (F), D-5 (G) or -16 h and +8 h (H), mice were administered 2 µg of murine IFN- λ 2 or PBS by intranasal route. Viral RNA levels from brain at 4 dpi (E: n = 7 per group, 2 experiments; F: n = 6-8 per group, 2 experiments; G: n = 6-8 per group, 2 experiments; H: n = 8 per group, 2 experiments). Bars indicate median values. Data were analyzed by Mann-Whitney test (***P* < 0.01 and ****P* < 0.001).



Figure S3. Cytokine responses following IFN- λ treatment and SARS-CoV-2 infection, Related to Figure 2. Eight-week-old female K18-hACE2 mice treated with 2 µg of murine IFN- λ 2 or PBS at -16 h by the intranasal route were challenged with 10³ FFU of WA1/2020 D614G. Cytokine levels in lung homogenates at 3 dpi (2 experiments, n = 7 per group except naïve n = 4). Data were analyzed by one-way ANOVA with Tukey's multiple comparison test (**P* < 0.05, ***P* < 0.01, ****P* < 0.001, and *****P* < 0.0001).



Figure S4. Cytokine induction following IFN- λ treatment and SARS-CoV-2 infection, Related to Figure 3. Six-week-old female 129S2 mice treated with two doses of 2 µg of murine IFN- λ 2 or PBS at -16 h and +8 h by the intranasal route were challenged with 10⁵ FFU of B.1.351 Beta variant. Cytokine levels in lung homogenates at 4 dpi (n = 7 per group except naïve n = 4, 2 experiments). Data analyzed by one-way ANOVA with Tukey's multiple comparison test (**P* < 0.05, ***P* < 0.01, ****P* < 0.001 and *****P* < 0.0001).



Figure S5. Heatmaps of RNA-seq data, Related to Figure 4. Heatmaps of selected significantly upregulated or downregulated gene sets corresponding with IFN- λ 2 treatment identified through GO analysis. Genes shown in each pathway are the union of the differentially expressed genes (DEGs) enriched in D+1 group or D+3 group versus control group (n = 4 per group). Columns represent sample groups and rows indicate genes.



Figure S6. Flow cytometric gating strategy and staining of lung cells, Related to Figure 5. (A) Six-week-old male and female C57BL/6 mice were inoculated with 10⁵ FFU of B.1.351 Beta variant. *Ifnl2* and *Ifnl3* mRNA levels from lungs were measured at indicated days after infection by qRT-PCR (n = 6 per group, 2 experiments) (ND, not detectable, qRT-PCR Ct value >40). (B-E) For lung tissues, cells were gated on single, live, CD45⁺ and CD45⁻ cells. Alveolar macrophages (AM) were identified as CD45⁺ SiglecF^{hi} CD11c^{hi} cells, dendritic cells (DC) were identified as CD45⁺ SiglecF⁻ CD11c⁺ MHCII⁺ cells (B). B and T cells were identified as CD45⁺ CD19⁺ cells and CD45⁺ CD3⁺ cells, respectively (C). Neutrophils (N ϕ) and epithelial cells (EC) were identified as CD45⁺ CD11b⁺Ly6G⁺ cells and CD45⁻ CD326⁺ cells, respectively (D). Monocytes (Mo) were identified as CD45⁺ CD11b⁺Ly6G^{hi} cells (E). (F) Localization of EGFP and SARS-CoV-2 nucleocapsid protein (NP) in the lungs of WT C57BL/6 (non-reporter, negative control) mice at 2 dpi. Frozen sections were stained for GFP (green), NP (magenta), and Hoechst (blue). Scale bar, 50 µm. (G) *Ifnl2* and *Ifnl3* mRNA levels from lungs of six-week-old male and female naïve, uninfected WT, *Mavs^{-/-}*, *cGas^{-/-}* and *Myd88^{-/-}* C57BL/6 mice were measured by qRT-PCR (n = 6-8 per group, 2 experiments) (ND, not detectable, qRT-PCR Ct value > 40).



Figure S7. Flow cytometry analysis of peripheral blood and lungs from neutrophil-depleted or bone marrow chimeric mice, Related to Figure 6. (A) Experimental scheme of neutrophil deletion in 129S2 mice. (B) (*Left*) Representative flow cytometry plots of peripheral blood at D+4 following intraperitoneal injection of a depleting anti-Ly6G mAb (1A8) or isotype control mAb. (*Right*) Frequency of mature neutrophils (CD11b⁺Ly6B⁺Ly6G⁺Ly6C^{int}) in blood are shown after antibody depletion. (C) Frequency of mature neutrophils (CD11b⁺Ly6B⁺Ly6G⁺Ly6G⁺Ly6G⁺Ly6C^{int}) in lungs are shown after antibody depletion (n = 8 per group, 2 experiments). (D) Representative flow cytometry plots of peripheral blood at 10 weeks after irradiation and bone marrow cell transplantation of CD45.2 cells to CD45.1 recipient mice.