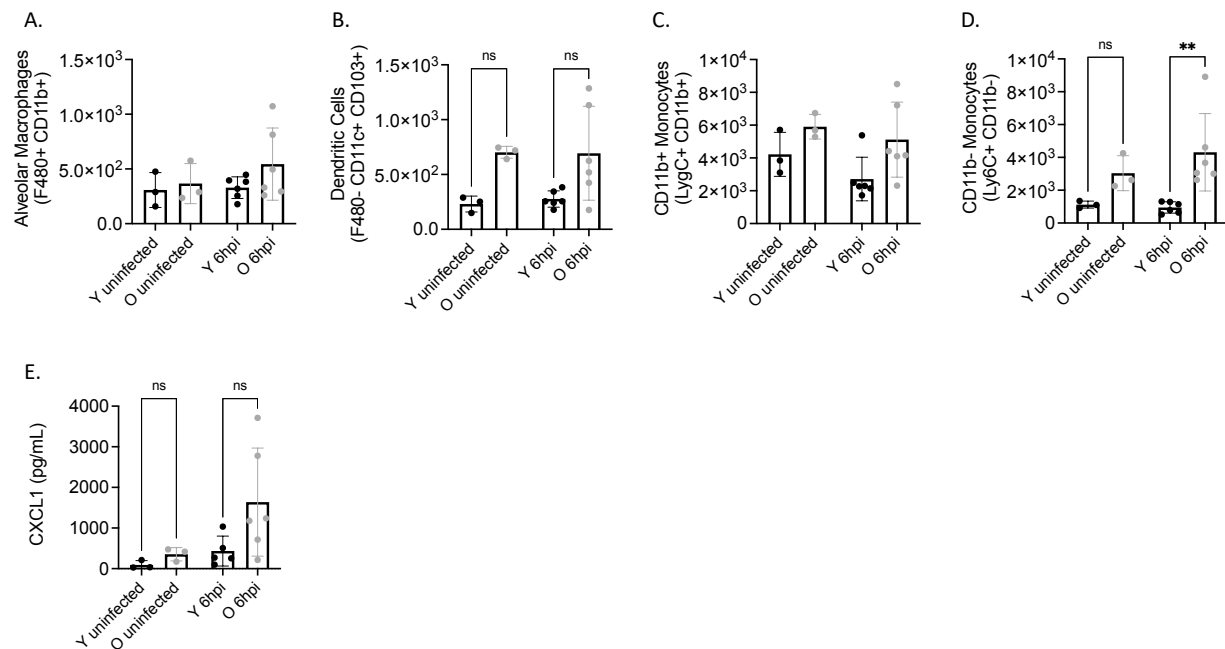


1 SUPPLEMENTAL MATERIAL



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3 **Supplemental Figure 1: Levels of innate immune cells and CXCL1 in young versus old**

4 **mice.** Young and old, male C57BL/6 mice were infected i.t with 5x10⁴ CFU *S. pneumoniae*

5 TIGR4 and at 6 hpi, lungs were harvested. Lungs were digested and stained for (A) Alveolar

6 Macrophages (F480+ CD11b+), (B) Dendritic Cells (F480- CD11c+ CD103+), (C) CD11b+

7 Monocytes (Ly6C+ CD11b+) and (D) CD11b- Monocytes (Ly6C+ CD11b-). (E) lungs were

8 homogenized and CXCL1 levels were measured by ELISA. Data are pooled from n=3 uninfected

9 and n=6 infected mice. Asterix indicates significance as determined by Kruskal-Wallis test, ns

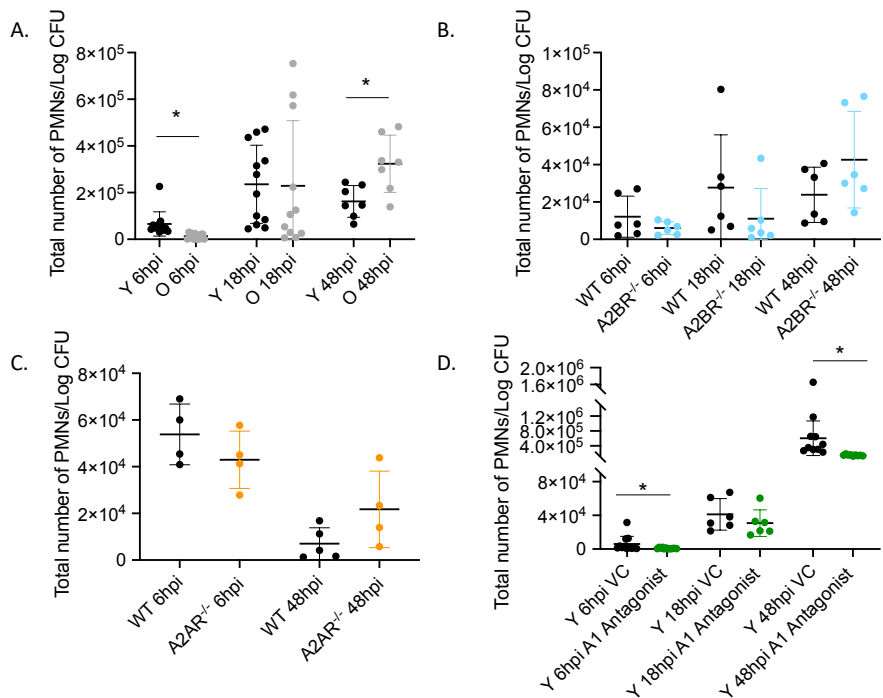
10 indicates not significant.

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Supplemental Figure 2: Total PMNs/Log CFU in Lung. (A) Young and old male C57BL/6 mice

(B), A2BR^{-/-} mice, (C) A2AR^{-/-} mice and (D) Young C57BL/6 mice treated with A1 inhibitor were

infected intra-tracheally (i.t.) *S. pneumoniae* TIGR4. Lungs were harvested, digested, and stained

for PMNs and analyzed by flow cytometry at the indicated timepoints following infection. Lung

samples were plated on blood agar and CFU was enumerated at the indicated timepoint post

infection. (A-D) For each mouse, the number of PMNs in the lungs were divided by the total CFU

per lung. (A) Data are pooled from 6 separate experiments with a total of n=7 mice at 48-hour time

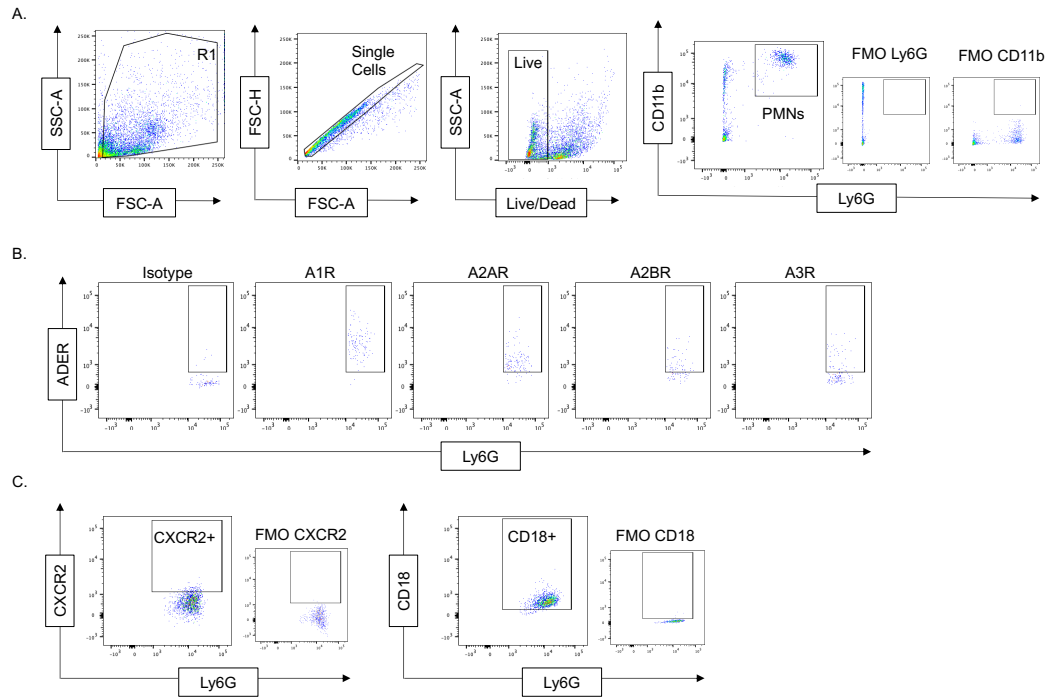
point and n=12 mice at all other time points. (B) Data are pooled from 4 separate experiments with

n=6-8 young mice per group. (C) Data are pooled from 2 experiments with n=11-12 young mice

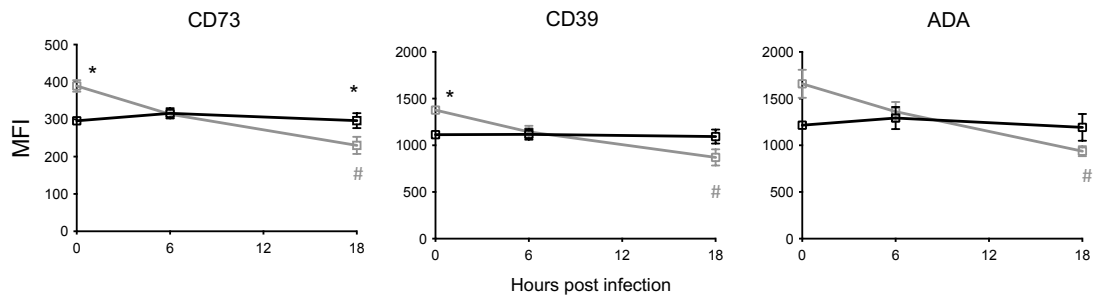
per group. (D) Data are pooled from 3 separate experiments with n=6-12 mice per group. Asterix

indicates significance as determined by Kruskal Wallis followed by Dunn's Multiple comparison

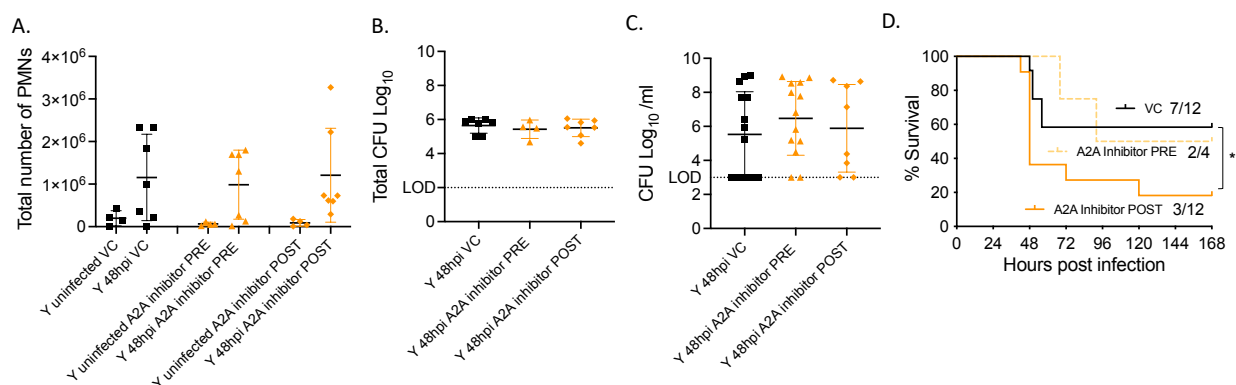
test.



Supplemental Figure 3: Gating Strategy. Lungs and blood were harvested, and receptor expression was determined by flow cytometry. Representative gating strategies are shown. (A) single cell, live, PMNs (Ly6G⁺, CD11b⁺) were gated on and (B) expression of the different adenosine receptors as well as (C) expression of CXCR2 and CD18 on PMNs was determined. Abbreviations: SSC-A= side scatter peak area, FSC-A= forward scatter peak area, FSC-H= forward scatter peak height, FMO= fluorescence minus one.

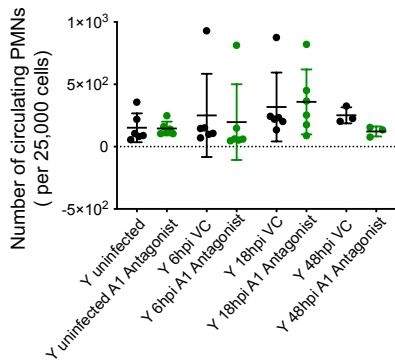


Supplemental Figure 4: EAD pathway enzyme expression in lungs. Young and old male C57BL/6 mice were infected i.t. with 5×10^4 CFU *S. pneumoniae* TIGR4. The Lungs were harvested, and lung digests were stained for the indicated adenosine producing and degrading enzymes. MFI on the surface of PMNs was determined over time using flow cytometry. Asterix indicates significance between young vs old at the indicated timepoint and # indicates significance for each group from its uninfected baseline as determined by Kruskal Wallis followed by Dunn's multiple comparisons test. Data are pooled from 2 separate experiments with n=12-14 mice per group.



Supplemental Figure 5: Signaling via A2A receptor does not affect PMN pulmonary numbers

or host resistance to infection. Young, male mice were treated i.p. with specific A2A receptor inhibitor or vehicle control (VC) either 18 hours prior to infection (PRE) or 18 hpi (POST). Mice were then infected i.t. with 5×10^4 CFU *S. pneumoniae* TIGR4. (A) Lungs were harvested 48 hpi, digested, and stained for PMNs. Samples were analyzed by flow cytometry. (B) Lung homogenates and (C) blood samples were plated on blood agar and bacterial CFU was enumerated. (A-C) Data are pooled from 3 separate experiments with (A) n=4-8 per group, (B) n=4-6 per group, (C) n=8-14 mice per group. (D) Following treatment and infection mice were monitored for survival. Asterix indicates significance as determined by Log-Rank test. Fractions indicate the number of surviving mice over the number of total mice per group. Data are pooled from two separate experiments.



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68 **Supplemental Figure 6: A1 receptor inhibition does not affect the number of circulating**

69 **PMNs.** Young, male C57/B6 mice were treated i.p. with specific A1 inhibitor and infected i.t with

70 5×10^4 CFU *S. pneumoniae* TIGR4. At indicated time points blood was harvested and stained for

71 PMNs (Ly6g+, CD11b+) and analyzed by flow cytometry. Data are pooled from 3 separate

72 experiments with n=6-12 mice per group.

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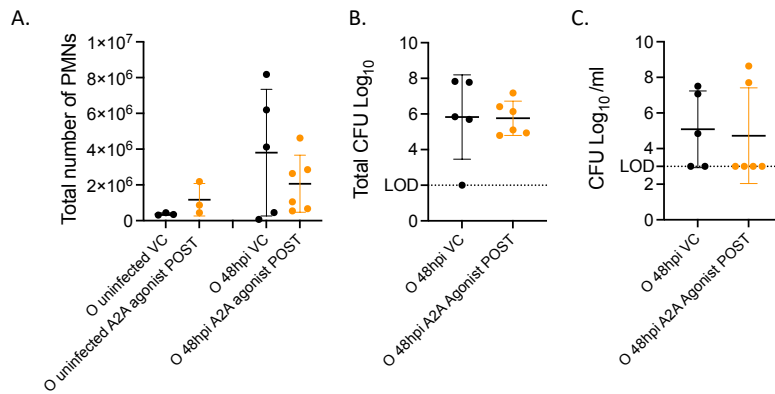
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Supplemental Figure 7: A2A receptor signaling does not affect late PMN pulmonary migration in aged hosts. Old mice were infected at 5×10^4 CFU with *S. pneumoniae* TIGR4 and 18 hpi treated i.p. with specific A2A agonist. 48 hpi lungs were harvested and total number of PMNs were determined by flow cytometry (A). Lungs were homogenized (A) and blood was collected (C) and plated on blood agar plates and CFU was enumerated. Data are pooled from n=3-6 mice per group.