

Supplementary Fig. 1. The IAV proteins released from H3-nanovax formulations remain structurally intact. Proteins liberated from the three H3-nanovax formulations and the input purified HA, NA, NP, or M1 proteins (Sino) were examined by SDS PAGE under reducing conditions. Gels were transferred onto a PVDF membrane, membranes blocked in 5% milk for 1 hr, washed, and incubated with primary antibody against their indicated protein overnight at 4°C. Membranes were then washed and incubated with goat anti-rabbit horseradish peroxidase (HRP) as well as ECL DualVue[™] Western Blotting Marker S-Protein-HRP to reveal molecular weight markers for 1-2 hrs, washed, and then reveled with an ECL substrate for 5-10 min. Images were obtained on a ChemiDoc MP Imager (Bio-Rad Laboratories). SDS-Page gels against hemagglutinin (HA: antibody 86001-RM01) (a), neuraminidase (NA: antibody 40040-MM02) (b), nucleoprotein (NP: antibody MA5-42364) (c), and matrix protein 1 (M1: antibody PA5-32253) (d). The predicted MW: Molecular weight kDa (kilodaltons) is indicated.



Supplementary Fig. 2. Lymphocyte Gating Tree. Lymphocyte Gating Tree. C57BL/6 mice were vaccinated i.n. with 500 µg of H3-nanovax encapsulating CpG + either 1% each HA + NP, M1 + NA, or NA + NP or left unvaccinated (naïve). At 45- or 50- days post vaccination lungs and draining lymph nodes were harvested and stained for expression of markers of interest. B and T cells were segregated based on CD19, CD4, and CD8α expression. B cells within the lung vs those in the circulation were further identified based on CD45.2i.v.Ab⁻. B cells within the lung were further subset into FAS⁺ B cells (CD45.2i.vAb⁻CD19⁺Fas⁺) and Germinal Center (GC) B cells (CD45i.vAb⁻CD19⁺ Fas⁺GL7⁺). Fas⁺GL7⁻ B cells were further subset into resident memory B cells (B_{RM}; CD45i.vAb⁻CD19⁺ Fas⁺GL7⁻CD38⁺CD69⁺IgM⁻CXCR3⁺). T cell populations in CD19⁻ cells were gated based on CD4 or CD8α expression and then further subset into antigen-experienced populations based on CD11a, CD49d, CD44 expression (AgExp CD4; CD4⁺CD11a^{hi}CD49d⁺ and AgExp CD8; CD11a^{hi}CD44⁺. Airway resident T cells were identified on the basis of CD11a expression (airway CD4/CD8 T cells; CD11a^{lo}CD44⁺). Antigen-experienced lung-resident memory T_{RM} populations of CD4 and CD8 T cells were determined on the basis of CD45.2i.v.Ab⁻, CD69 and CD103 expression.



Supplementary Fig. 3. Germinal Center B cells and B_{RM} in the lungs following H3-nanovax. Supplemental Figure 2. Germinal Center B cells and B_{RM} in the lungs following H3-nanovax. C57BL/6 mice were vaccinated i.n. with 500 µg of H3-nanovax (HA/NP; Pink), (NA/NP; Blue), (NA/M1; Purple), or left unvaccinated (naïve). Representative flow plots (a, b) and enumeration (c, d) of lung resident germinal center (GC) B cells (CD45i.vAb⁻CD19⁺ Fas⁺GL7⁺, shown are CD45i.vAb⁻CD19⁺ gated cells) (a, c), and resident memory B cells (B_{RM}) (CD45i.vAb⁻CD19⁺ Fas⁺GL7⁻CD38⁺CD69⁺lgM⁻CXCR3⁺, shown are CD45i.vAb⁻CD19⁺ Fas⁺GL7⁻CD38⁺CD69⁺lgM⁻CXCR3⁺, shown are CD45i.vAb⁻CD19⁺ Fas⁺GL7⁻CD38⁺CD69⁺ cells) (b, d) at 45 or 50 days post vaccination. Error bars, mean ± s.e.m. Data representative of one (NA/NP, NA/M1: n= 5 mice/group) or two (Naïve, HA/NP: n= 4 mice/group) independent experiments. * P< 0.05, ** P< 0.01 (One-way ANOVA with Tukey's multiple comparisons test).



Supplementary Fig. 4. H3-nanovax formulations induce lymph noderesident Germinal Center B cells responses. C57BL/6 mice were vaccinated i.n. with 500 µg of H3-nanovax (HA/NP; Pink), (NA/NP; Blue), (NA/M1; Purple), or left unvaccinated (naïve). Representative flow plots (a) and enumeration (b) of lymph node resident germinal center (GC) B cells (Fas⁺GL7⁺) at 45- or 50-days post vaccination. Shown are CD45i.vAb⁻CD19⁺ gated cells. Error bars, mean ± s.e.m. Data representative of two (NA/M1: n= 3-4 mice/group) or three (Naïve, HA/NP, and NA/NP: n= 4-5 mice/group) independent experiments. * P< 0.05, ** P< 0.01 (One-way ANOVA with Tukey's multiple comparisons test).



Supplementary Fig. 5. CpG-nanovax does not induce significant immunity relative to unvaccinated mice. C57BL/6 mice were vaccinated as described in Fig. 2. 32 days following prime vaccination, lungs were harvested and GC and B_{RM} (a, b) B cell responses were enumerated. Total numbers of antigen experienced lung (CD11a^{hi}CD44⁺) (c, f), airway (CD11a^{lo}CD44⁺) (d, g), and CD4 (CD69⁺CD103⁻, e) and CD8 (CD69⁺CD103⁺,h) Trm T cells were also enumerated. Error bars, mean ± s.e.m. Data representative of one n= 3-4 mice/group) independent experiment. * P< 0.05, (One-way ANOVA with Tukey's multiple comparisons test).



Supplementary Fig. 6. H3-nanovax encapsulating IAV proteins induces payload targeted systemic and lung-local IAV-specific antibody responses. C57BL/6 mice were vaccinated with H3-nanovax encapsulating CpG + either HA/NP (Pink), NA/NP (Blue), NA/M1 (Purple), or left unvaccinated (naïve). At 45 days post-vaccination, serum and BAL were collected and total IAV H3N2 A/Hong Kong/1/68 (i.e., virion specific) serum and BAL IgG (a, b) and BAL IgA (c) were quantified via ELISA. Error bars, mean \pm s.e.m. Data representative of two independent experiments with n = 4 mice/group. (a-c) HA/NP vs. naïve: * P< 0.05, ** P< 0.01, **** P< 0.001; NA/NP vs. naïve: # P< 0.05, ## P< 0.01, #### P< 0.001; NA/M1 vs. naive: \ddagger P< 0.05, \ddagger P< 0.01, ### P< 0.001; NA/NP vs. naïve: \ddagger P< 0.001, #### P< 0.001, NA/NP vs. NA/M1: @ P< 0.05, @@ P< 0.01, @@@ P< 0.001, @@@@ P< 0.0001; NA/NP vs. NA/M1: \ddagger P< 0.05, \ddagger P< 0.01, \ddagger P< 0.001, "the P< 0.0001, "the P< 0.001, "the P< 0.00



Supplementary Fig. 7. IAV-nanovax induced antibody is specific to the viral protein component rather than the His-tag. C57BL/6 mice were vaccinated as described in Fig. 2. At 45- or 50- days post vaccination, serum and BAL were collected and ELISA and total anti-His-SARS-CoV-2 S1 subunit specific IgG was quantified to determine whether His-tags on the proteins utilized in H3-nanovax formulations elicited His-reactive antibody. Error bars, mean \pm s.e.m. Dotted line = 0. Data representative of two independent experiments with n = 4 mice/group. Two-way ANOVA with Tukey's multiple comparison test



CX3CR1 – BV711

Supplementary Fig. 8. H3-nanovax formulation generated antigen-experienced CD103⁺CD69⁺ CD8 T cells exhibit a uniform T_{RM} phenotype relative to CD103⁻ and circulating counterparts. C57BL/6 mice were vaccinated as described in Figure 2. 32 days following prime vaccination, lungs were harvested, and the phenotype of vaccine generated CD103⁺CD69⁺ (HA/NP: Pink, NA/NP: Blue, NA/M1: Purple) and CD103⁻CD69⁺ (Gray) antigen-experienced CD8 T cells in the lung interstitium was determined using established T_{RM} markers Eomesodermin (Eomes) (a), CD49a (b), CXCR3 (c), and CX3CR1 (d). Circulating antigen-experienced CD8 T cells (i.e. CD45.2i.v.Ab⁺: Orange) served as a control. Data representative of one (n= 4-5 mice/group) independent experiment.



Supplementary Fig. 9. H3-nanovax formulations induce lymph noderesident T cell responses. C57BL/6 mice were vaccinated (H3-nanovax (HA/NP; Pink), (NA/NP; Blue), (NA/M1; Purple), or left unvaccinated (naïve)) as described in Fig. 2. Representative flow plots (a, b) and enumeration (CD4: c; CD8: d) of antigen-experienced CD4 (c, CD11a^{hi}CD49d⁺) and CD8 (b, CD11a^{hi}CD44⁺) T cells enumerated at 45- or 50- days post vaccination. Shown are CD4⁺CD11a^{hi}CD49d⁺) (a, c) and CD8a⁺CD11a^{hi}CD44⁺) (b, d) gated cells. Error bars, mean ± s.e.m. Data representative of two (M1/NA: n= 3-4 mice/group) or three (Naïve, HA/NP, and NA/NP: n= 4-5 mice/group) independent experiments. * P< 0.05, ** P< 0.01, *** P< 0.001 (One-way ANOVA with Tukey's multiple comparisons test).



Supplementary Fig. 10. H3-nanovax formulations generate vaccine specific CD8 T cell responses. C57BL/6 mice were vaccinated (CpG onlynanovax (light purple), H3-nanovax (HA/NP; Pink), (NA/NP; Blue), (NA/M1; Purple), or left unvaccinated (naïve)) as described in Fig. 2. Lung resident antigen-experienced CD8 (CD45i.vAb⁻CD8a⁺CD44⁺) T cells specific for NP₃₆₆₋₃₇₄ and M1₁₂₈₋₁₃₅ derived from IAV A/Hong Kong/1/68 and A/Aichi/2/68 respectively were enumerated (a, b, d, e). Enumeration of dLN antigen experienced (CD11a⁺CD44⁺) CD8 T cells specific for NP₃₆₆₋₃₇₄ (c). Error bars, mean \pm s.e.m. Data representative of one n= 3-4 mice/group) independent experiment. * P< 0.05, (One-way ANOVA with Tukey's multiple comparisons test).