

## Supplemental Information

### **ARNAX is an ideal adjuvant for COVID-19 vaccines to enhance antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses and neutralizing antibody induction**

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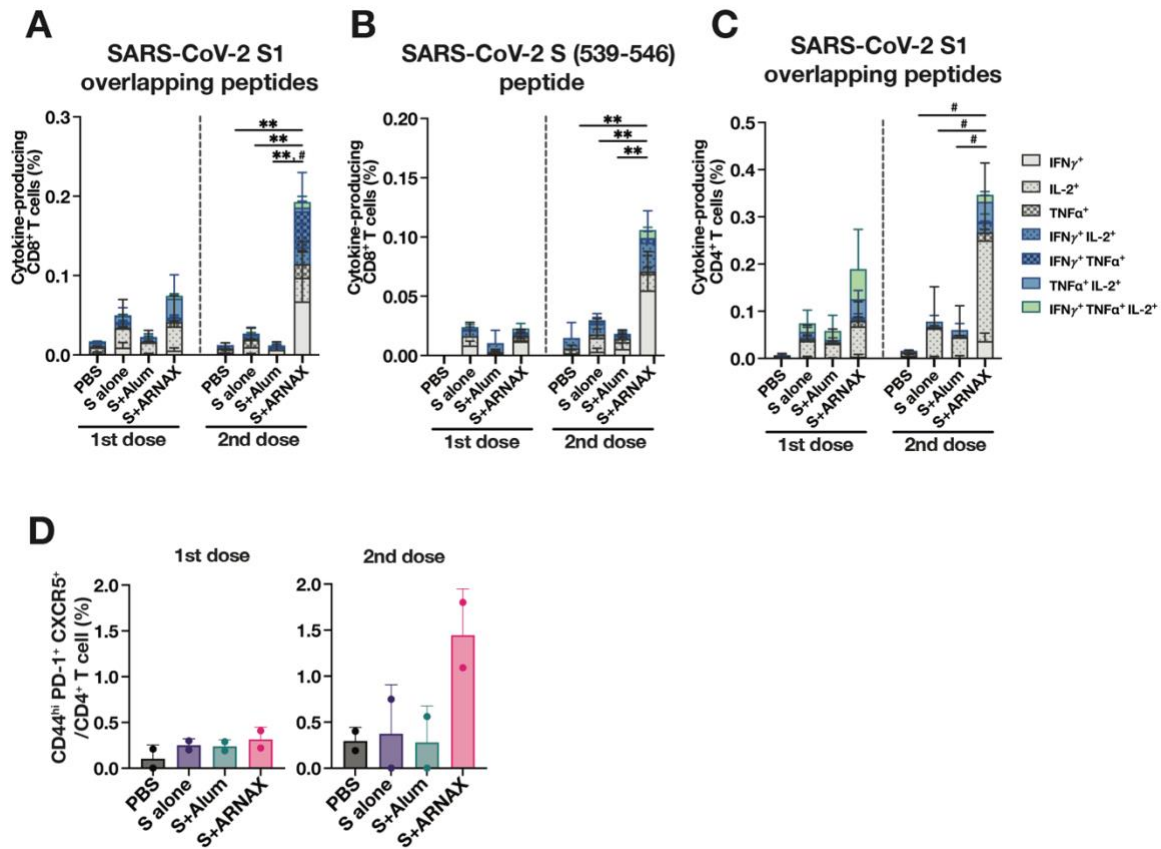
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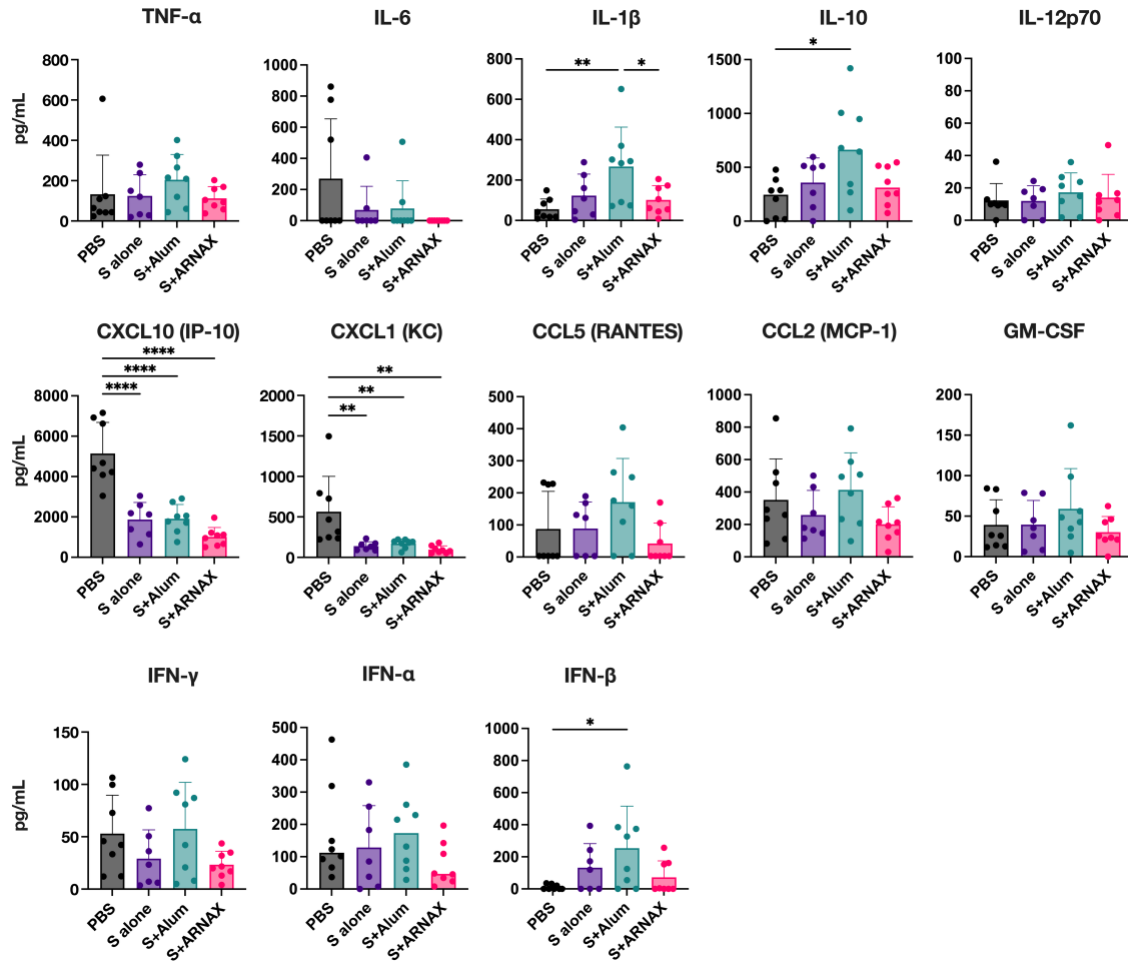
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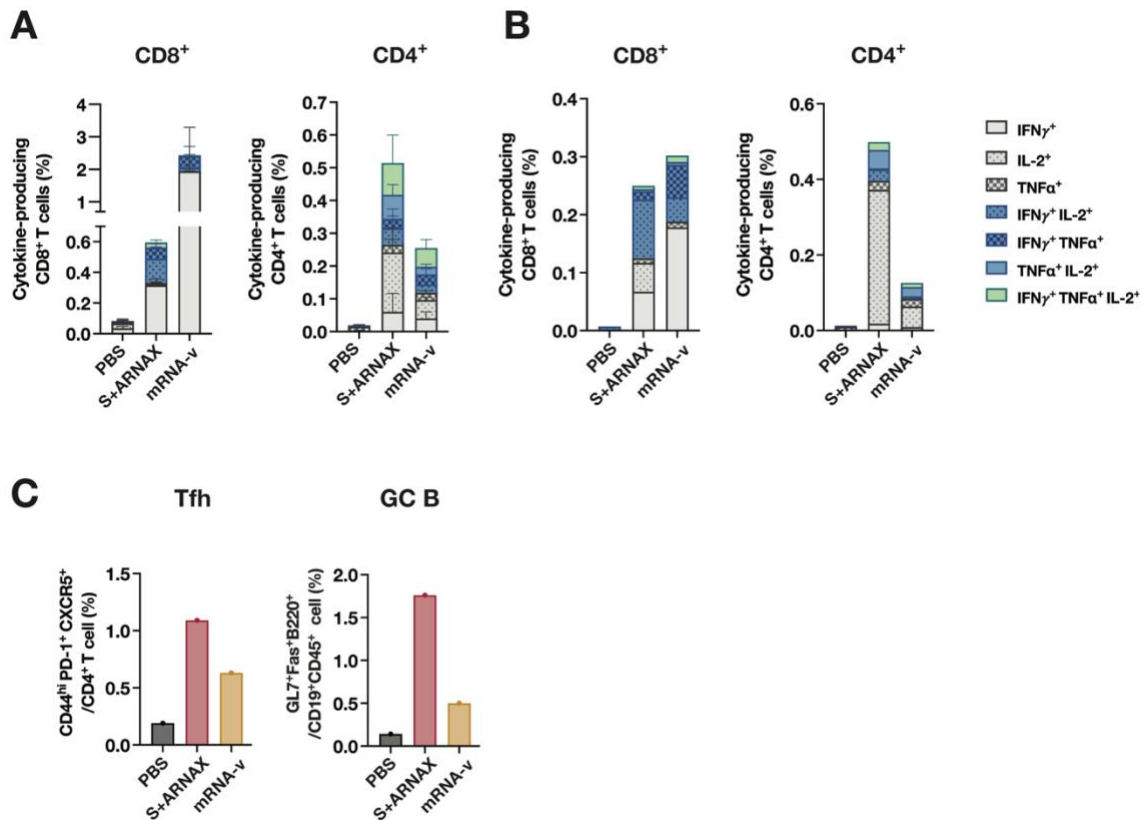
**Fig S1.** Analysis of cytokine producing T cells

Mice were immunized with each vaccine (n = 7-10). Ten days after the first dose or 4 days after the second dose of vaccine, lymph nodes were harvested. The lymph nodes from 3-4 mice for each experiment were pooled within the vaccine group as a single sample. The pooled lymph node cells were re-stimulated with overlapping peptides (A, C) or AA 539-546 peptide of S protein (B). The results of 3 independent experiments are combined and shown in A, B, and C. Percentages of cytokine-producing CD4<sup>+</sup> and CD8<sup>+</sup> T cells (IFN- $\gamma$ , IL-2, and TNF- $\alpha$ ) were analyzed by intracellular cytokine staining using flow cytometry. Individual cytokine patterns are represented by the following: light gray for IFN- $\gamma$ <sup>+</sup>, light gray with small dots for IL-2<sup>+</sup>, light gray with large dots for TNF- $\alpha$ <sup>+</sup>, light blue with small dot for IFN- $\gamma$ <sup>+</sup> IL-2<sup>+</sup>, light blue with large dot for IFN- $\gamma$ <sup>+</sup> TNF- $\alpha$ <sup>+</sup>, light blue for TNF- $\alpha$ <sup>+</sup> IL-2<sup>+</sup>, green for IFN- $\gamma$ <sup>+</sup> TNF- $\alpha$ <sup>+</sup> IL-2<sup>+</sup>. Each error bar represents mean  $\pm$  SD. Significance is calculated for total cytokine-producing cells and multiple cytokine-producing cells (colored bars) using Two-way ANOVA with Tukey's multiple comparisons test. For total cytokine-producing cells: \*\* $P$ <0.01, multiple cytokine-producing cells: # $P$ <0.05. The percentage of Tfh, defined as CD44<sup>hi</sup>, PD-1<sup>+</sup>, and CXCR5<sup>+</sup>, in CD4<sup>+</sup> T cells in lymph nodes was analyzed (D). The results of 2 independent experiments are combined and shown in D.



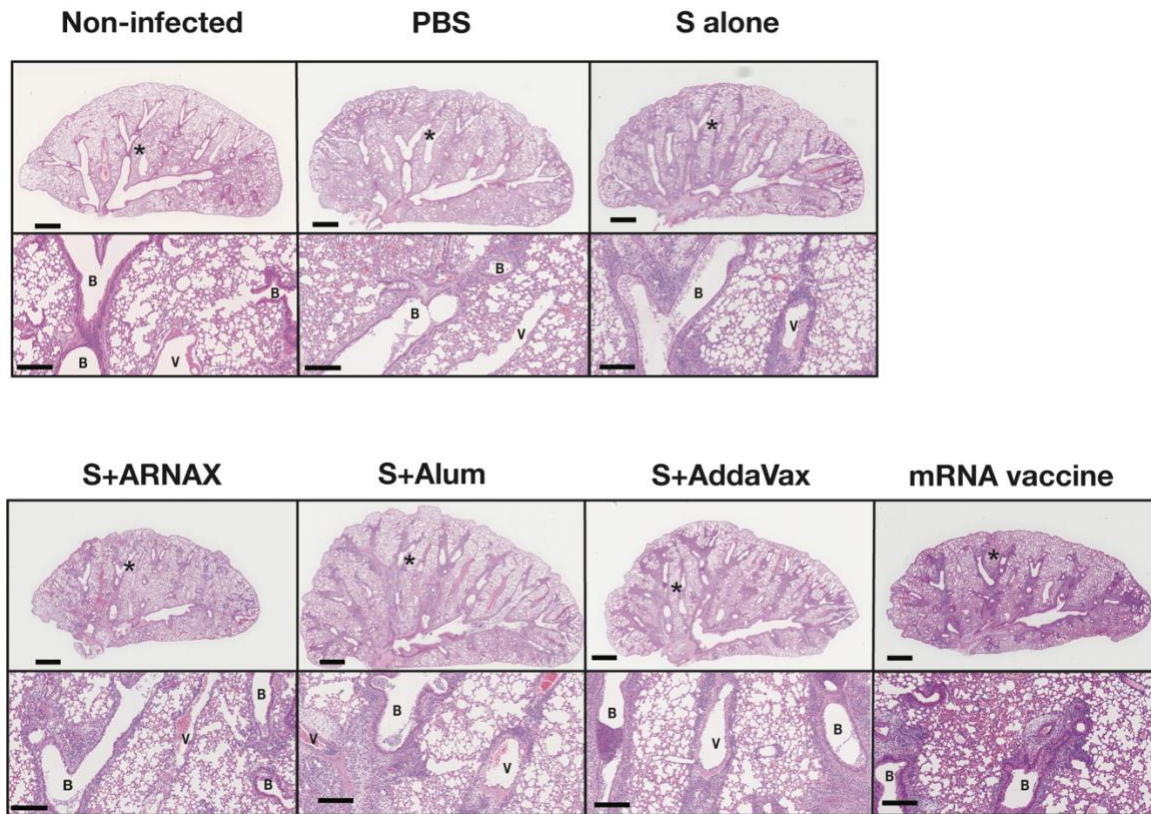
**Fig. S2.** Cytokine and chemokine levels in the sera at 3 days post-infection

Serum samples collected at 3 dpi were analyzed in a bead-based immunoassay to measure concentrations of cytokines and chemokines as indicated. Individual samples are shown as dots, horizontal lines indicate the median ( $n = 8$ ). Significance was calculated using One-way ANOVA with Tukey's multiple comparisons test. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.0001$ .



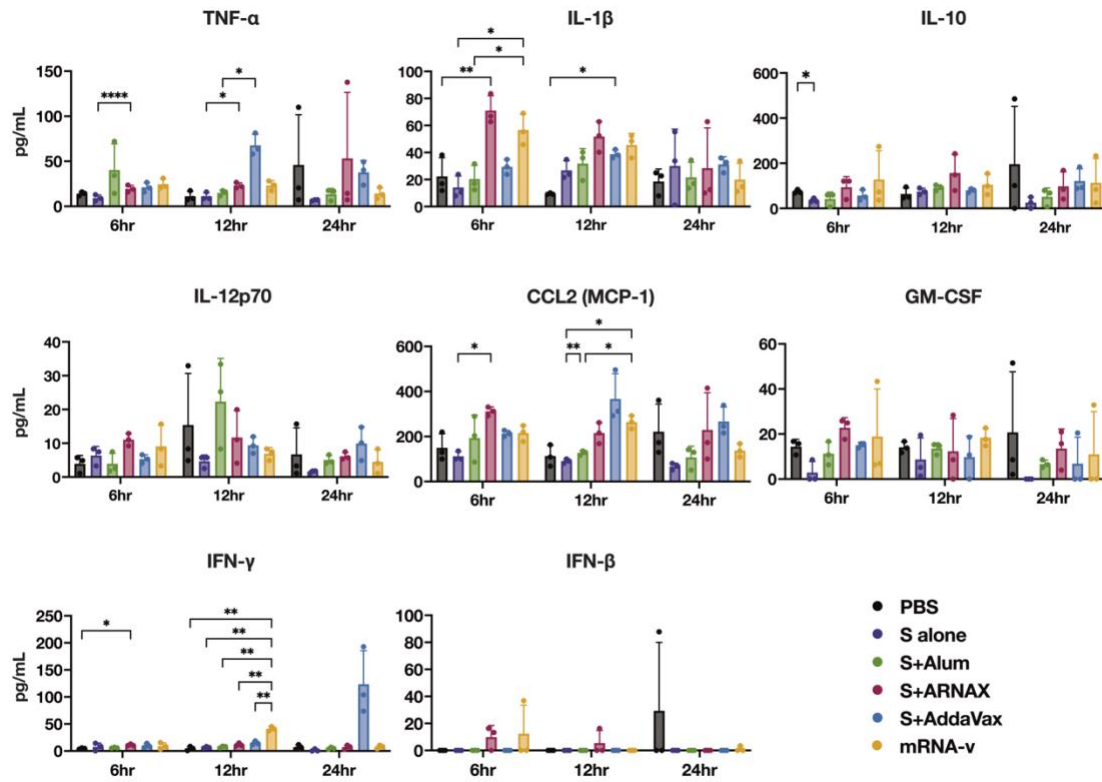
**Fig S3.** Comparison of T cell responses between S plus ARNAX and mRNA vaccines

Mice were immunized with each vaccine (n = 3). Four days after the second dose of vaccine, spleen and lymph nodes were harvested. The lymph nodes cells from 3-4 mice for each experiment were pooled within the vaccine group as a single sample. The splenocytes (A) and pooled lymph node cells (B) were re-stimulated with overlapping peptides of S protein. Percentages of cytokine-producing CD4<sup>+</sup> and CD8<sup>+</sup> T cells (IFN- $\gamma$ , IL-2, and TNF- $\alpha$ ) were analyzed by intracellular cytokine staining using flow cytometry. Individual cytokine patterns are represented by the following: light gray for IFN- $\gamma$ <sup>+</sup>, light gray with small dots for IL-2<sup>+</sup>, light gray with large dots for TNF- $\alpha$ <sup>+</sup>, light blue with small dot for IFN- $\gamma$ <sup>+</sup> IL-2<sup>+</sup>, light blue with large dot for IFN- $\gamma$ <sup>+</sup> TNF- $\alpha$ <sup>+</sup>, light blue for TNF- $\alpha$ <sup>+</sup> IL-2<sup>+</sup>, green for IFN- $\gamma$ <sup>+</sup> TNF- $\alpha$ <sup>+</sup> IL-2<sup>+</sup>. Each error bar represents mean  $\pm$  SD for splenocytes. The percentage of Tfh (defined as CD44<sup>hi</sup>, PD-1<sup>+</sup>, and CXCR5<sup>+</sup>) in CD4<sup>+</sup> T cells and GCB (defined as GL7<sup>+</sup>, Fas<sup>+</sup>, B220<sup>+</sup>) cells in CD19<sup>+</sup> CD45<sup>+</sup> B cells in lymph nodes was analyzed (C). Results are representative of at least two independent experiments.



**Fig S4. Histopathology of lung section at 3 dpi**

Mice were challenged with  $1 \times 10^5$  PFU of SARS-CoV-2 mouse-adapted strain 2 weeks after the second dose. The lung tissues were harvested at 3 dpi and stained with H&E. Representative pictures of each group ( $n = 5$ ) were shown. The asterisk of upper panel is indicated indicates an enlarged area (bottom panel). The scale bars indicated 1mm (upper panel) and 250  $\mu$ m. Bronchiole and vessel are marked as B and V respectively.



**Fig S5.** Cytokine and chemokine levels 6-24 hours after vaccination

Sera were collected from immunized mice at 6-24 hours after the first dose of vaccine, and concentrations of the indicated cytokines and chemokines were measured using a bead-based immunoassay ( $n = 3$ ). Individual samples are shown as dots, horizontal lines indicate the median. Significance was calculated using One-way ANOVA with Tukey's multiple comparisons test. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\*\* $P < 0.0001$ .

**Table S1** Cryo-EM data collection, refinement, and validation statistics

EMD-63071 spike protein	
EMDB entry	EMD-63071
Data collection and processing	
Magnification	130,000
Microscope	Krios G4
Voltage (kV)	300
Detector	Gatan K3
Energy filter	Gatan BioContinuum, 10 eV slit
Electric exposure (e-/Å)	52.85
Defocus range (μm)	−0.5 to −3.0
Pixel size (Å)	0.67
Data Processing Program	CryoSPARC (v.4.2.1)
Movies	3,500
Initial / Final particle images (no.)	14,448,075 / 31,127
Symmetry imposed	C1
Map resolution (Å)	3.67
FSC threshold	0.143