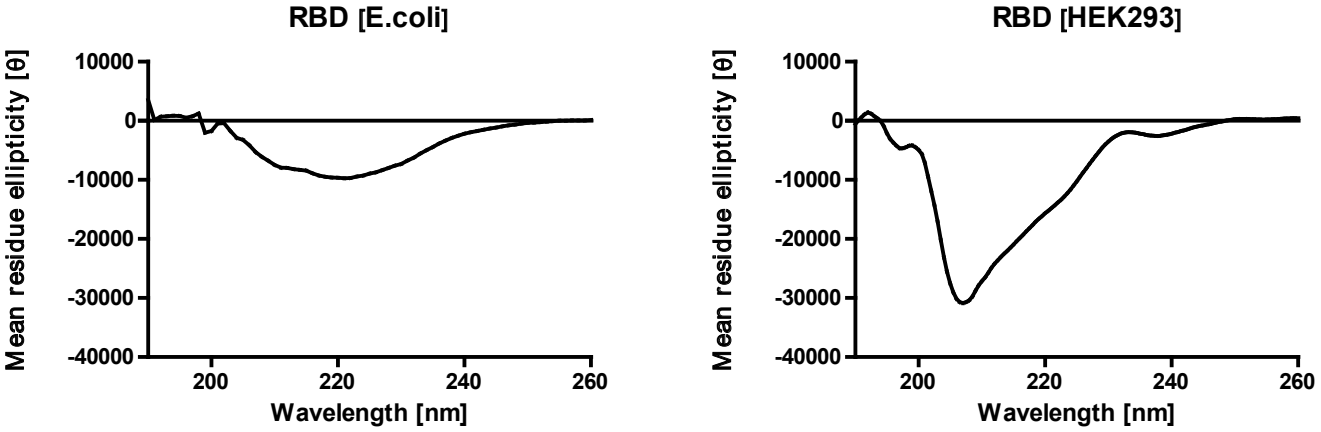
Figure S1. Circular dichroism analysis of (A) RBD expressed in E. coli or HEK293 cells[§] and (B) synthetic peptides 12 and 32. Molecular ellipticities (y-axes) at given wave lengths (x-axes) are shown.
[§]adapted from [39] (Gattinger et al. 2022, doi: 10.1111/all.15305).

(A)



(B)

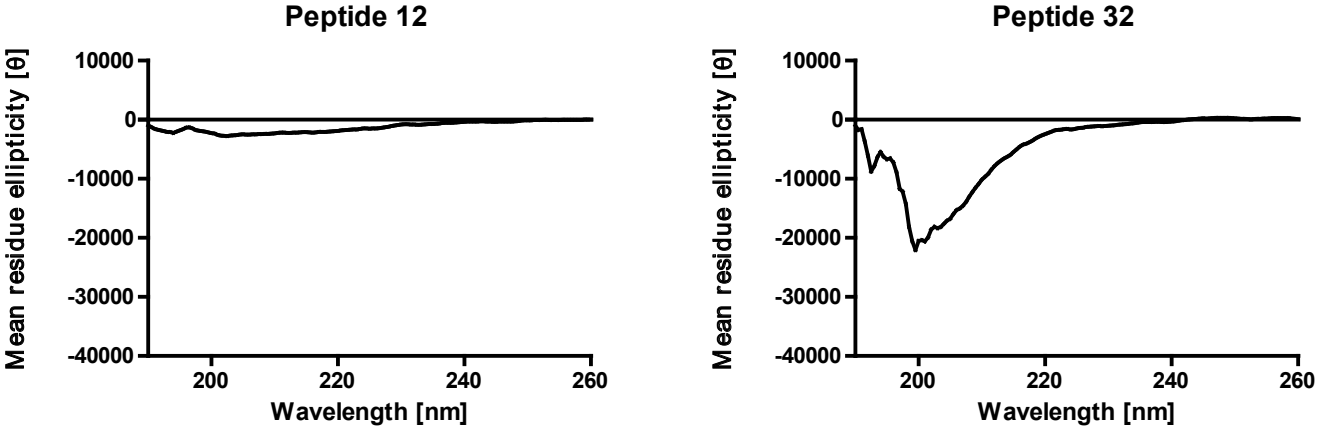


Figure S2. IgG antibody levels (y-axes: ISU) specific for RBD, S, S1 and S2 in naïve (colored in blue) and convalescent (colored in red) subjects at time point 0 and 3. The Friedman test was used for comparison between groups. Asterisks indicate significant difference between groups, *P < 0.05, ***P < 0.001, ns - not significant.

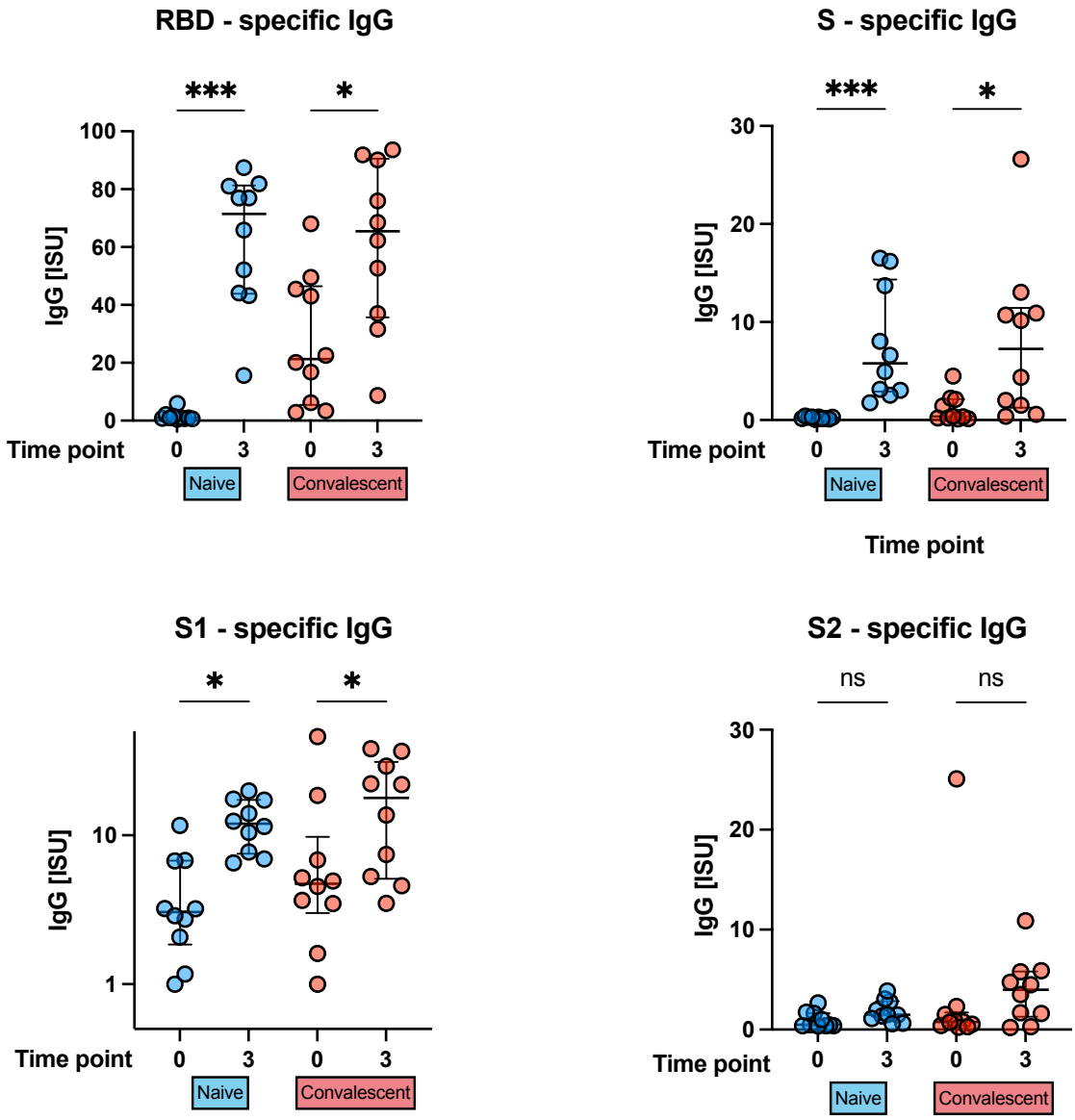


Figure S3. Comparison of IgG antibody levels (y-axes: ISU) specific for RBD, S-protein, S1 and S2 (x-axes) between naïve (colored in blue) and convalescent (colored in red) subjects at time point 3. The Mann-Whitney test was used for comparison between two groups. Asterisks indicate significant difference between groups, ns - not significant.

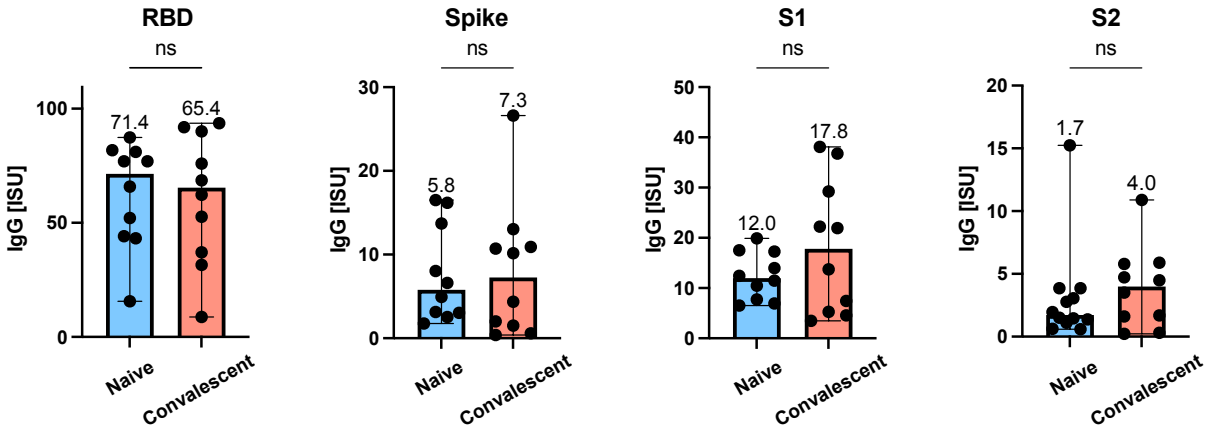


Figure S4. Changes of IgG antibody levels (y-axes: delta mean fluorescent intensities MFI) to S-derived peptides (x-axes) in naïve (lower part) and convalescent subjects (upper part) between time points 0 and 3.

