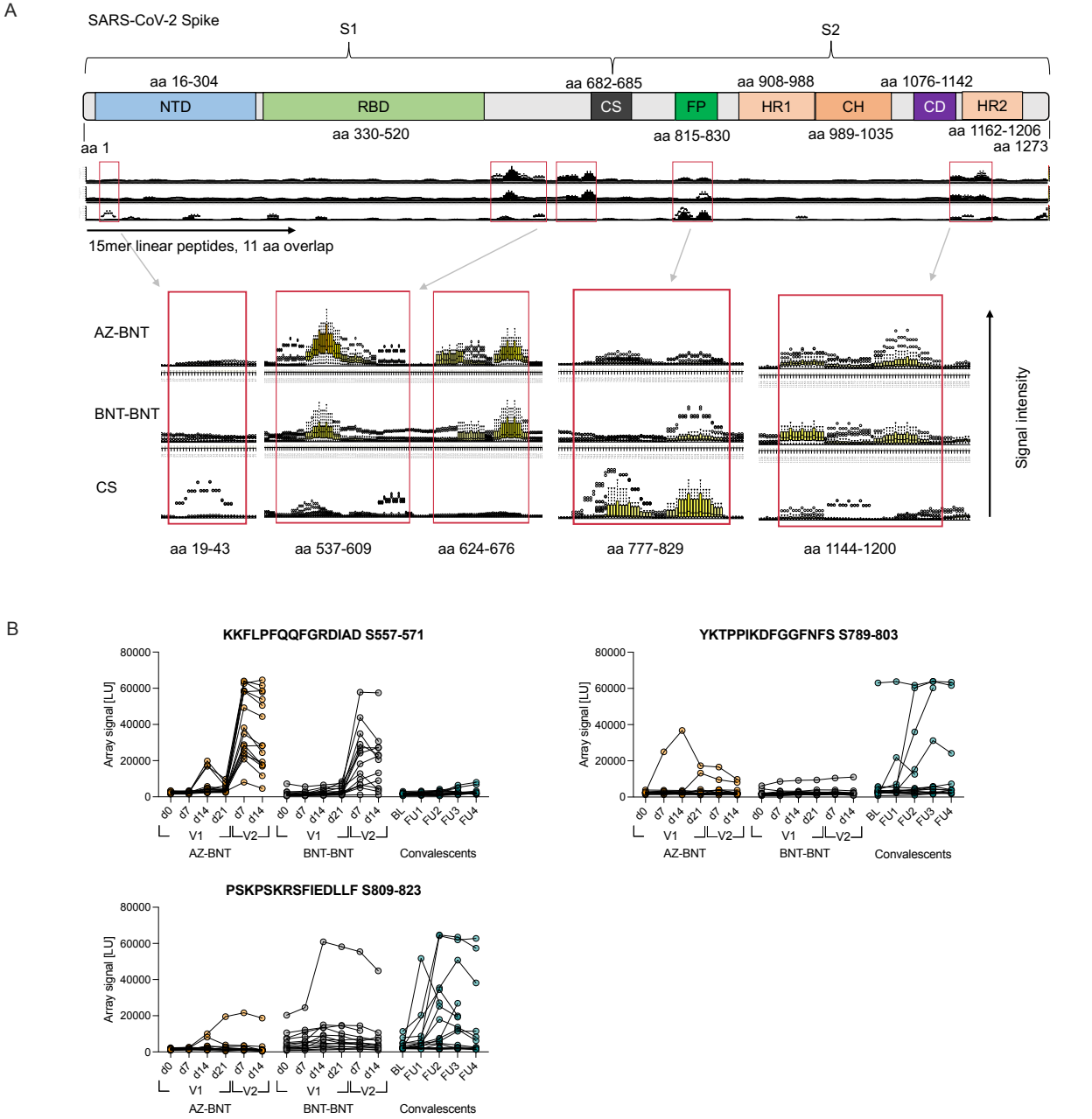

Supplementary Figure 3: Humoral spike epitope coverage following SARS-CoV-2 vaccination and infection. (A) Schematic overview of the spike glycoprotein structure (top row) and corresponding humoral responses to a peptide microarray with linear 15mers overlapping in 11 aa. Signal intensity per amino acid is shown for AZ-BNT (n=16, top row), BNT-BNT (n=15, middle) and convalescents (CS, n=17, bottom row) at day 7 post secondary vaccination and follow-up 3, respectively. NTD, N-terminal domain; RBD, receptor binding domain; CS, cleavage site; FP, fusion peptide; HR, heptad region; CH, central helix; CD, connector domain. Modified from Wrapp et al., Science 367, 1260-1263 (2020). **(B)** Signal from peptide microarray for selected peptides following AZ-BNT vaccination, BNT-BNT vaccination or infection (convalescents (CS)). BL = baseline; FU = follow up.