



S1 Fig. Correlation of IgG recognition of peptide motifs with overlapping amino acids. The fold enrichment in IgG with specificity for the different pairs of amino acid motifs (respective x and Y-axis) are shown along with the Spearman rank correlation ( $r_s$ ) value. Each symbol represents a different COVID-19 mRNA vaccine recipient. Trend lines were fitted by linear regression. Spearman rank correlation ( $r_s$ ) is displayed.