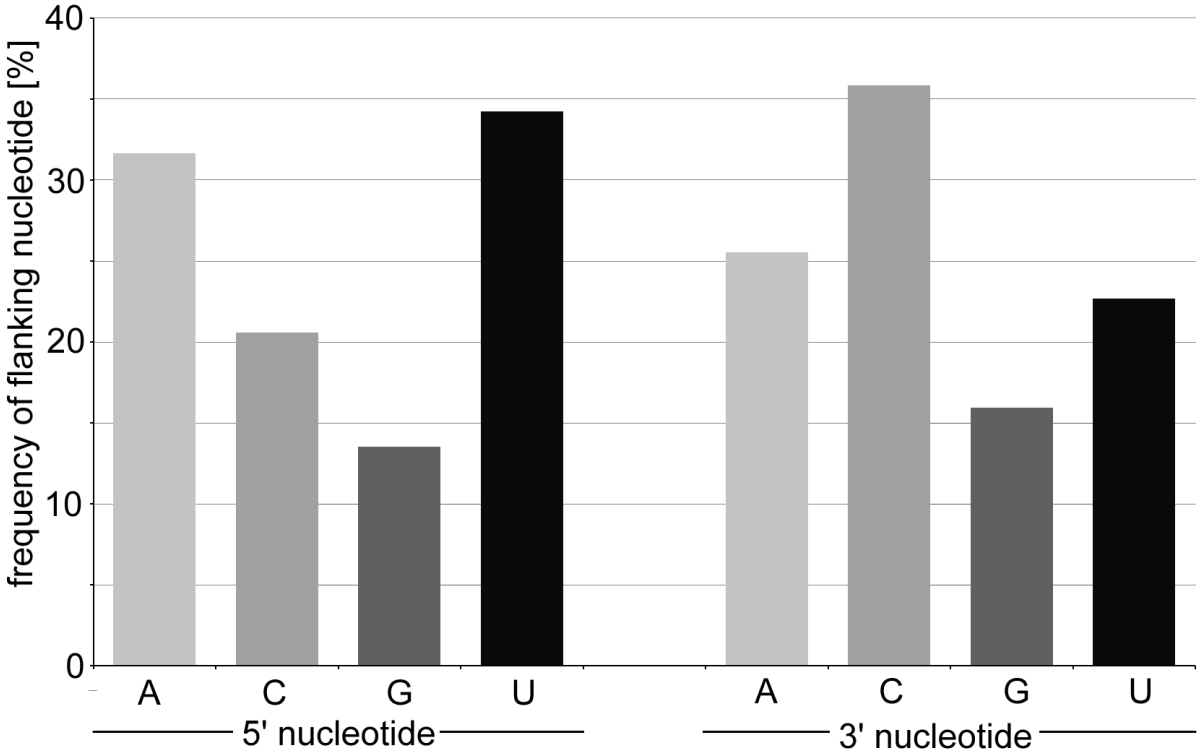Figure S5. Base frequency in motifs flanking the C>U substitutions. The source data are in **Table S8**. Nucleotide frequencies were normalized to base (A,C,G,U) distribution in the SARS-CoV-2 genome.