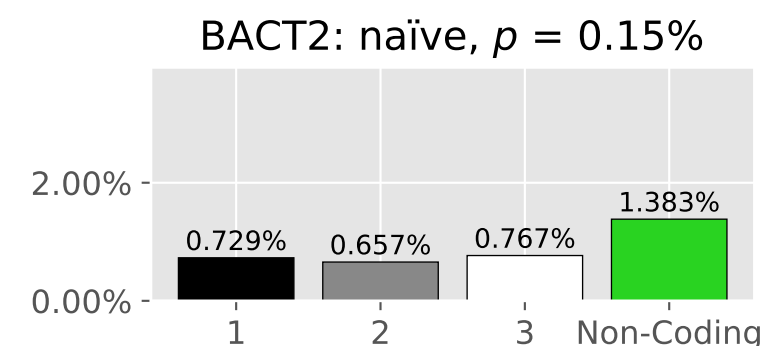
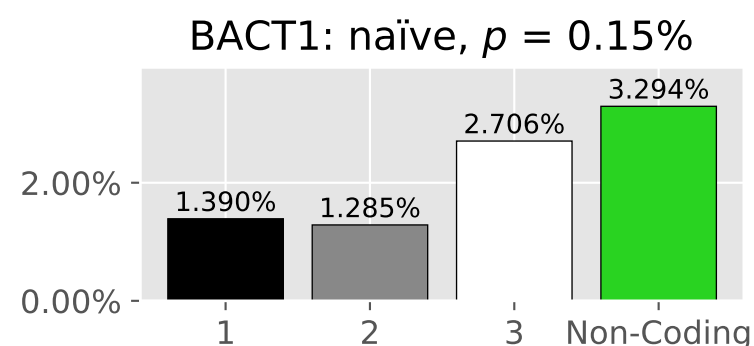
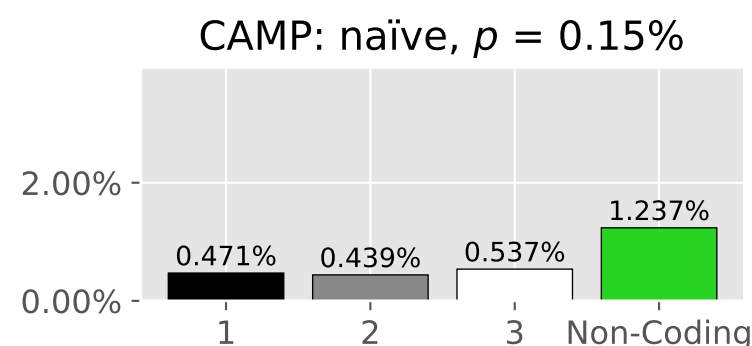
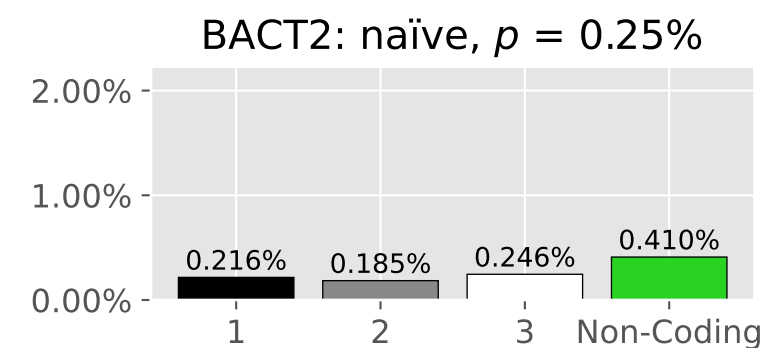
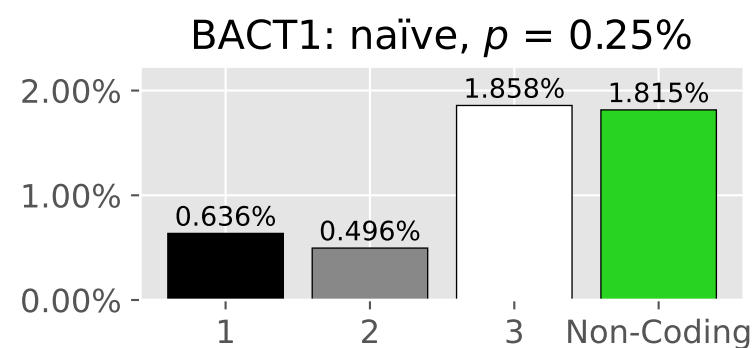
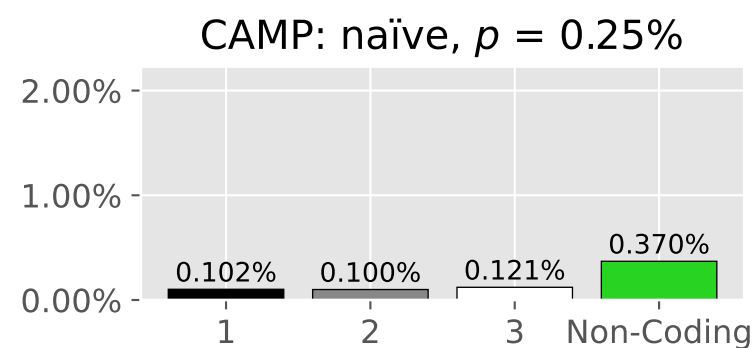
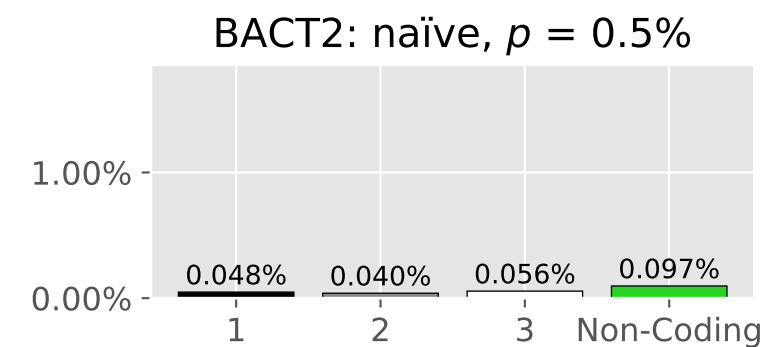
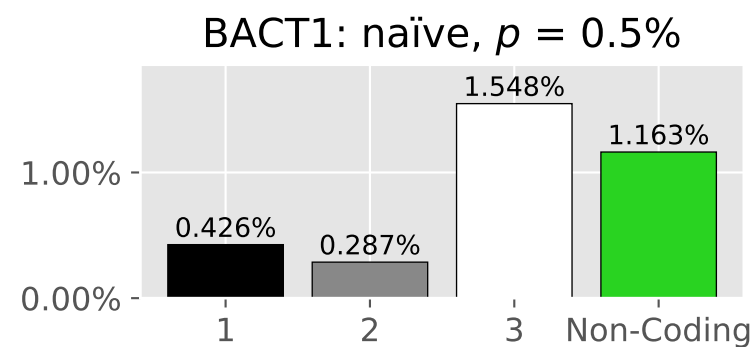
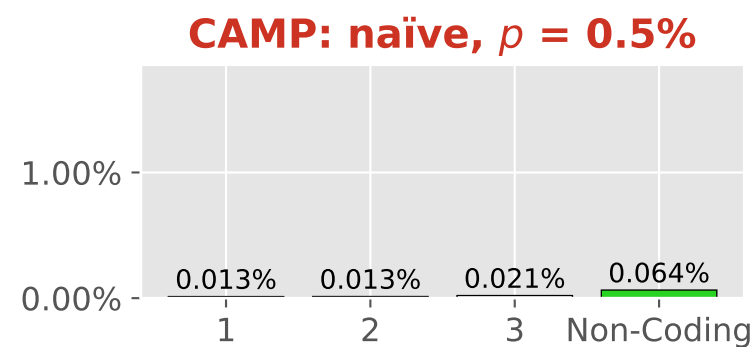
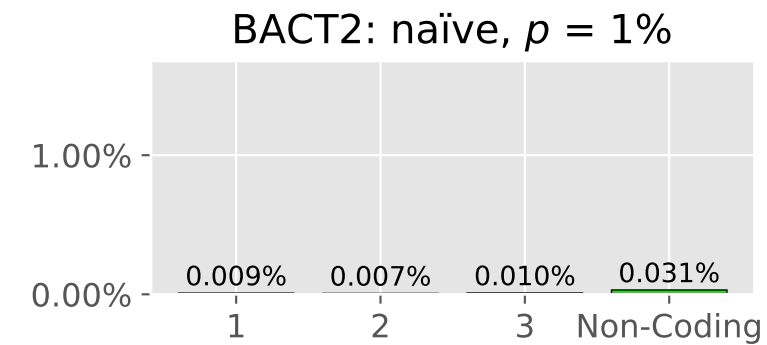
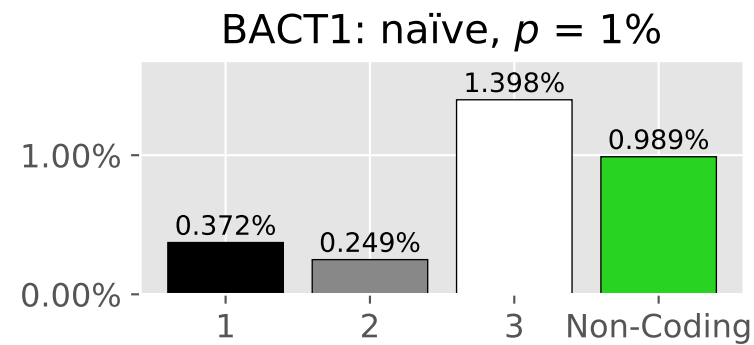
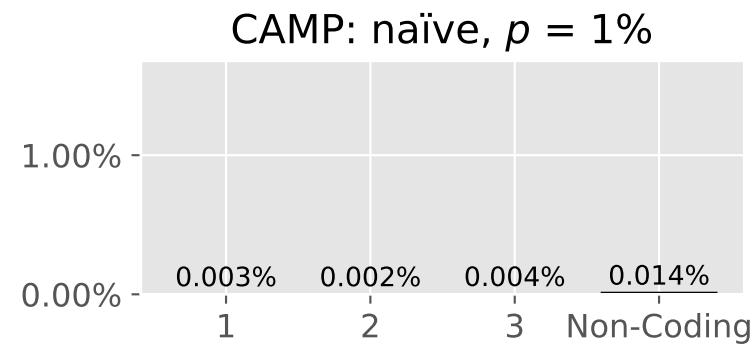
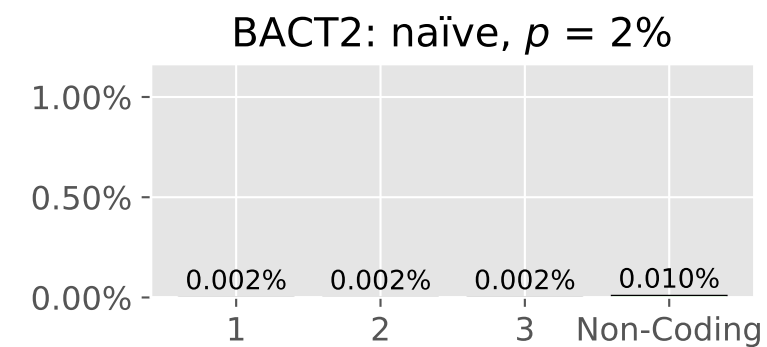
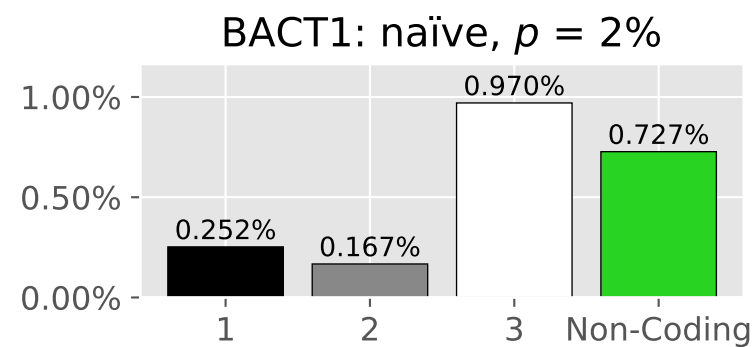
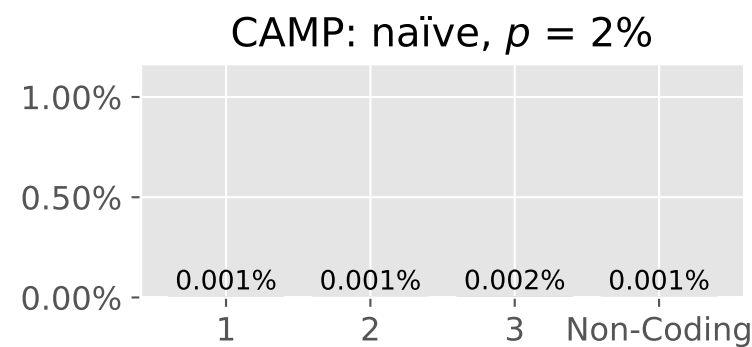
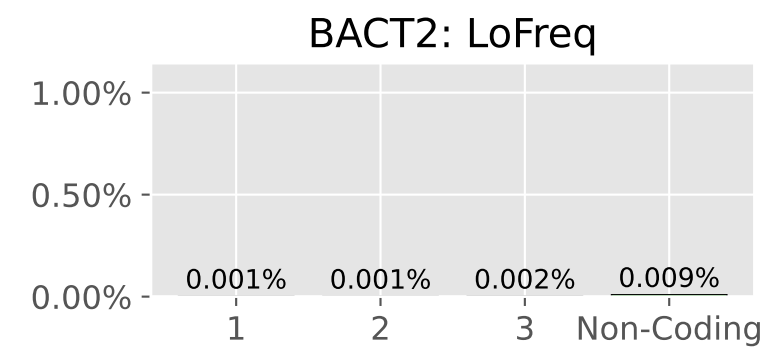
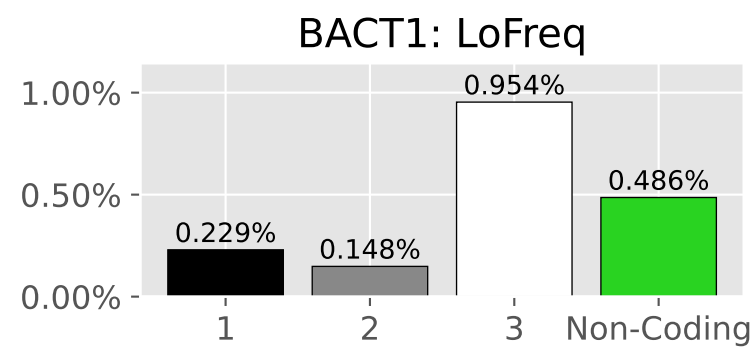
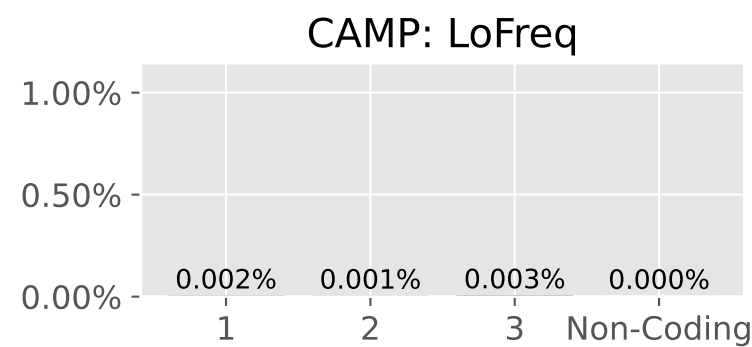


Rare mutation frequencies across codon positions (each bar normalized by total number of positions in that category)

Number of Mutated Positions
Number of Positions



Codon Position