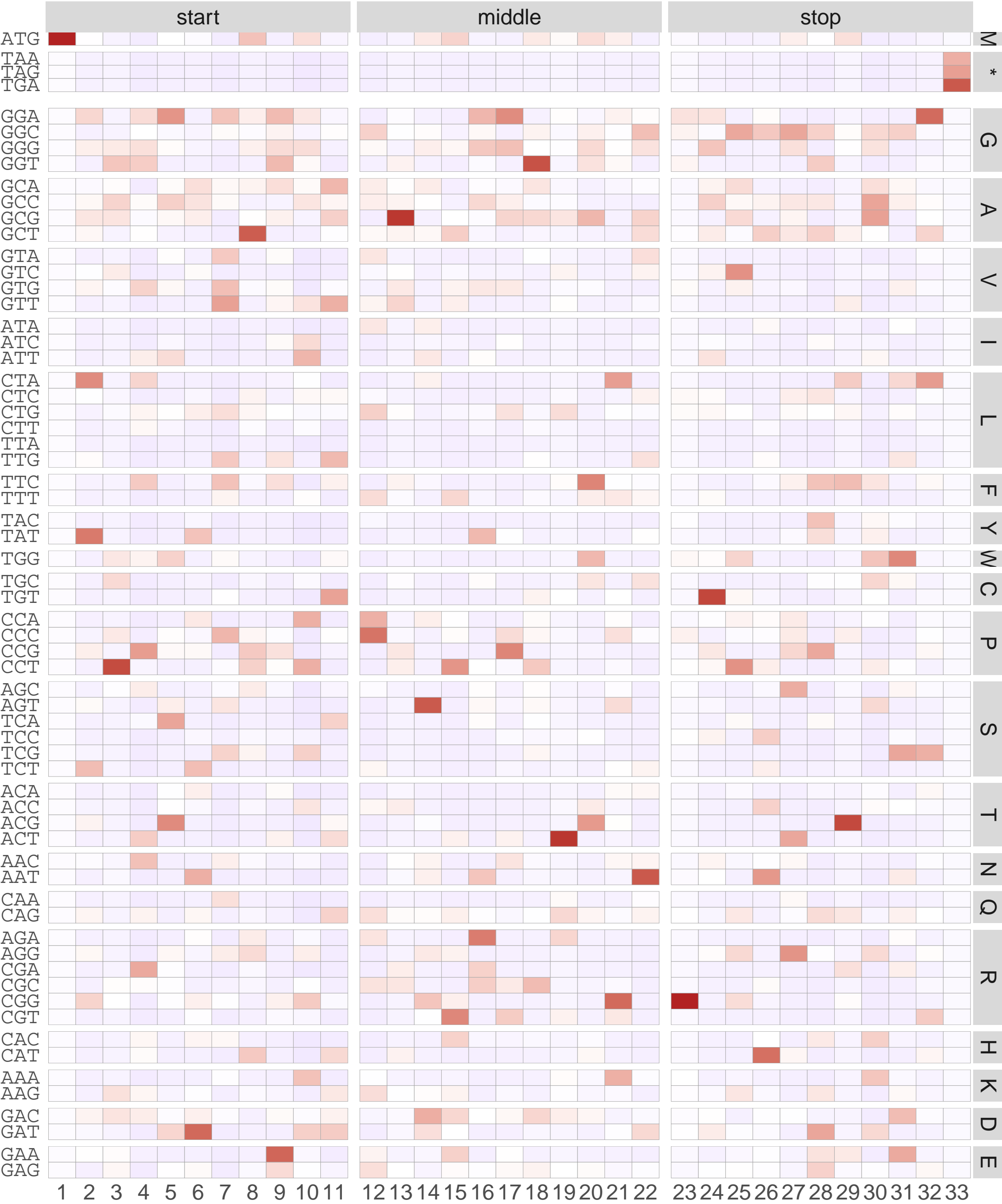


NA P-sites per codon(scale: zscore) 0 2 4 6



P-sites per codon(scale: zscore) 0 2 4 6 NA