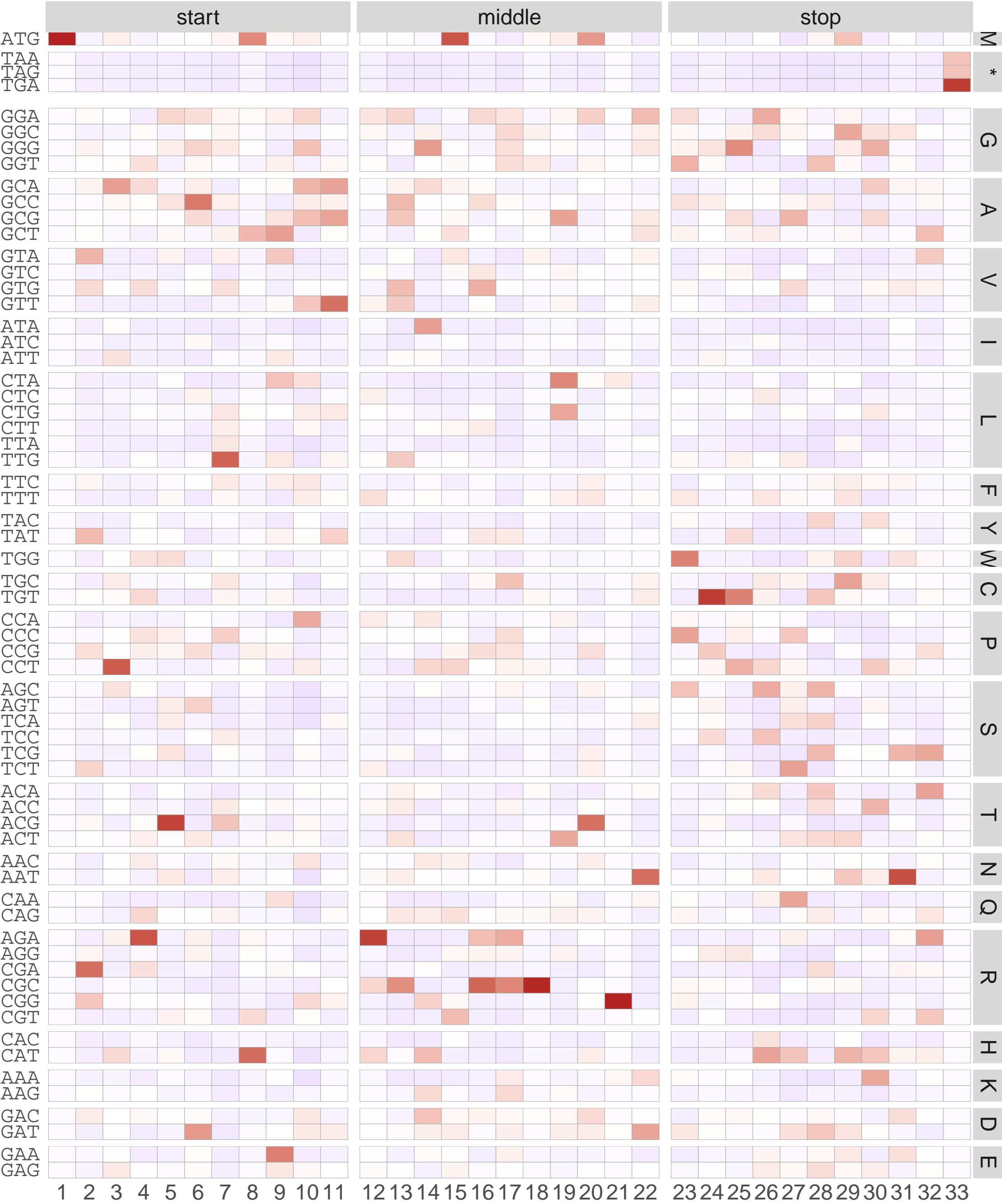


NA P-sites per codon(scale: zscore) 0 2 4 6



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