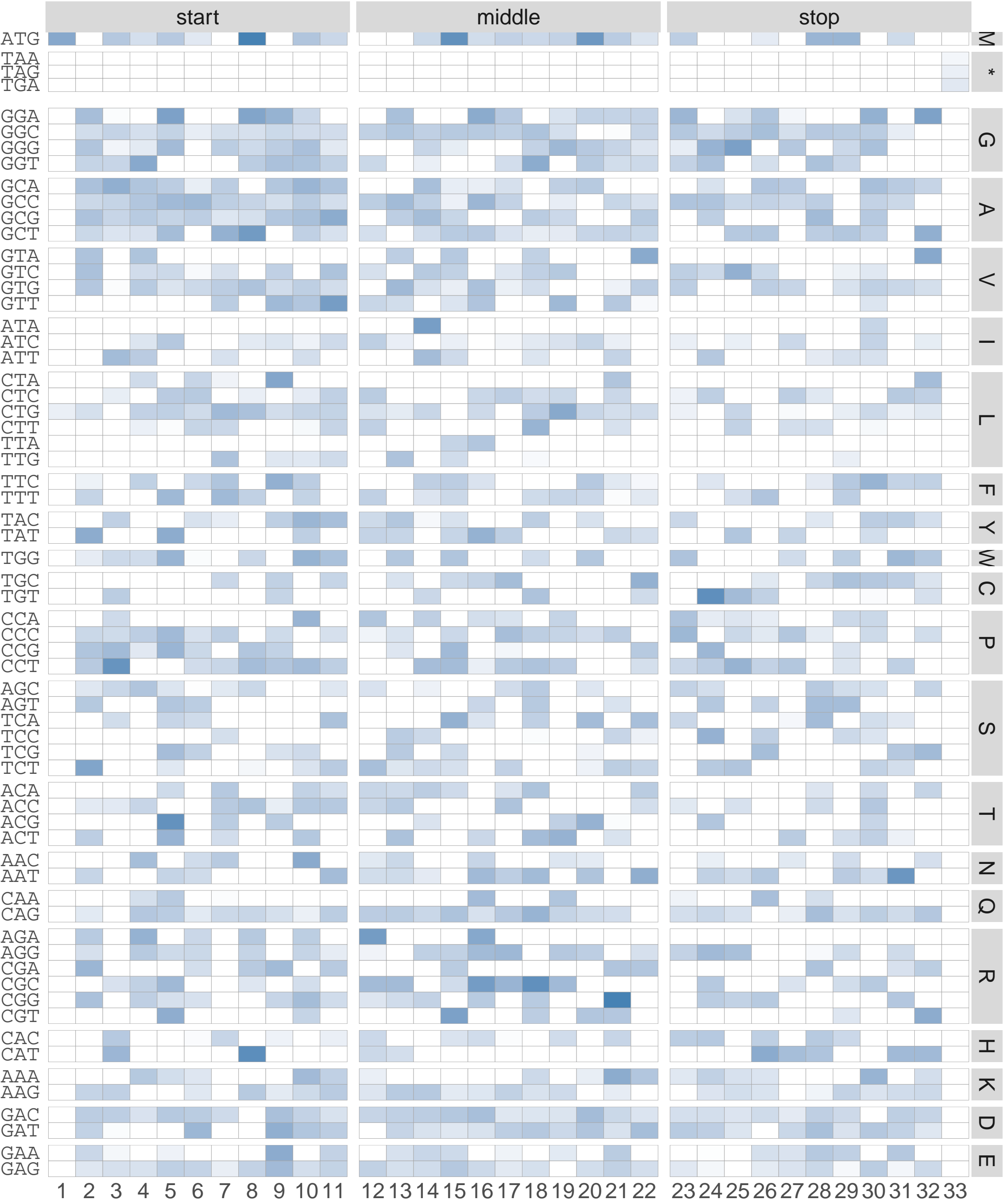


P-sites per codon(scale: log2) 0 2 4 6 NA



position (codon resolution)

P-sites per codon(scale: log2) 0 2 4 6 NA