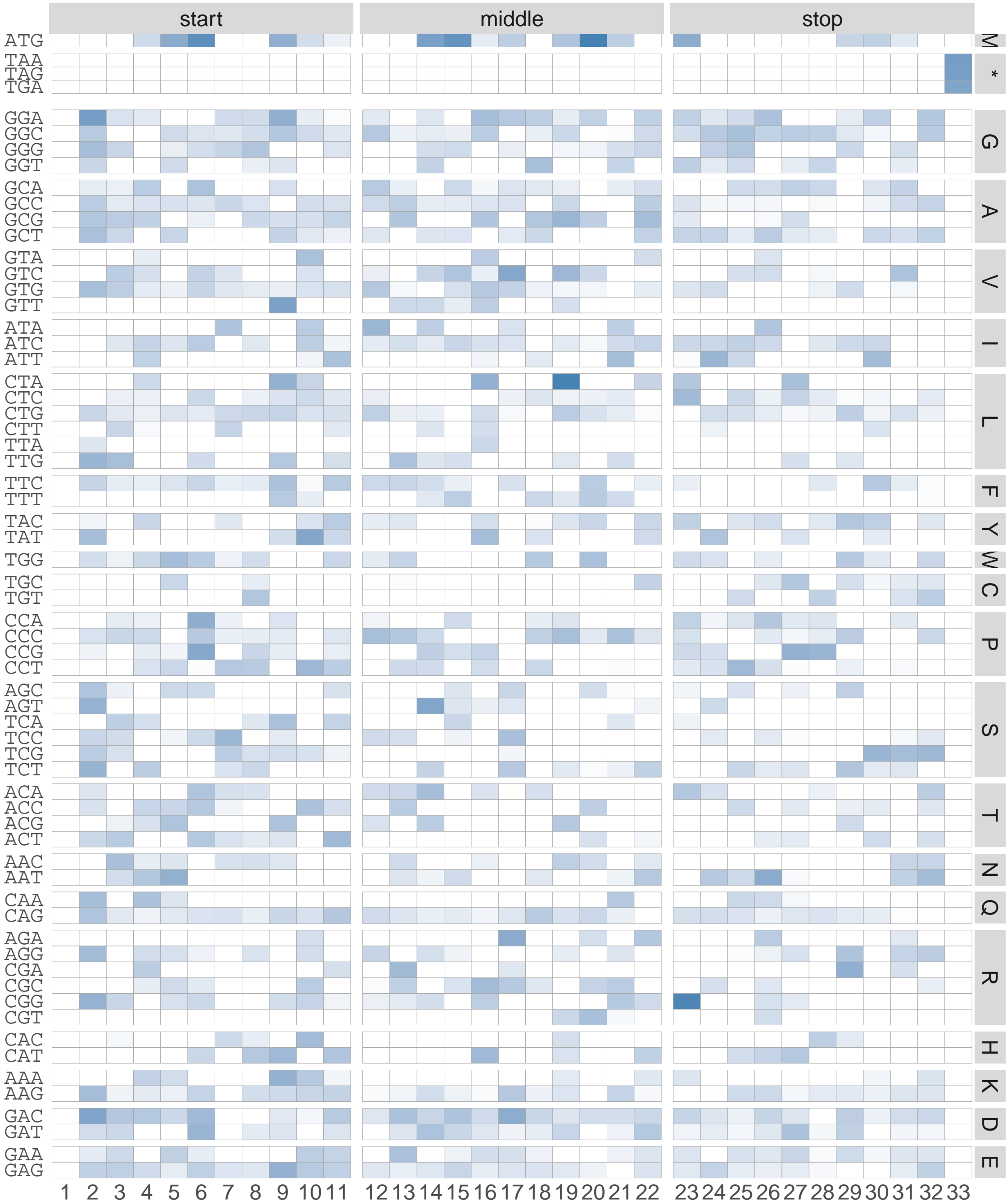


NA A-sites per codon(scale: log2) 0 1 2 3



position (codon resolution)

NA

A-sites per codon(scale: log2)

0 2 4 6