E−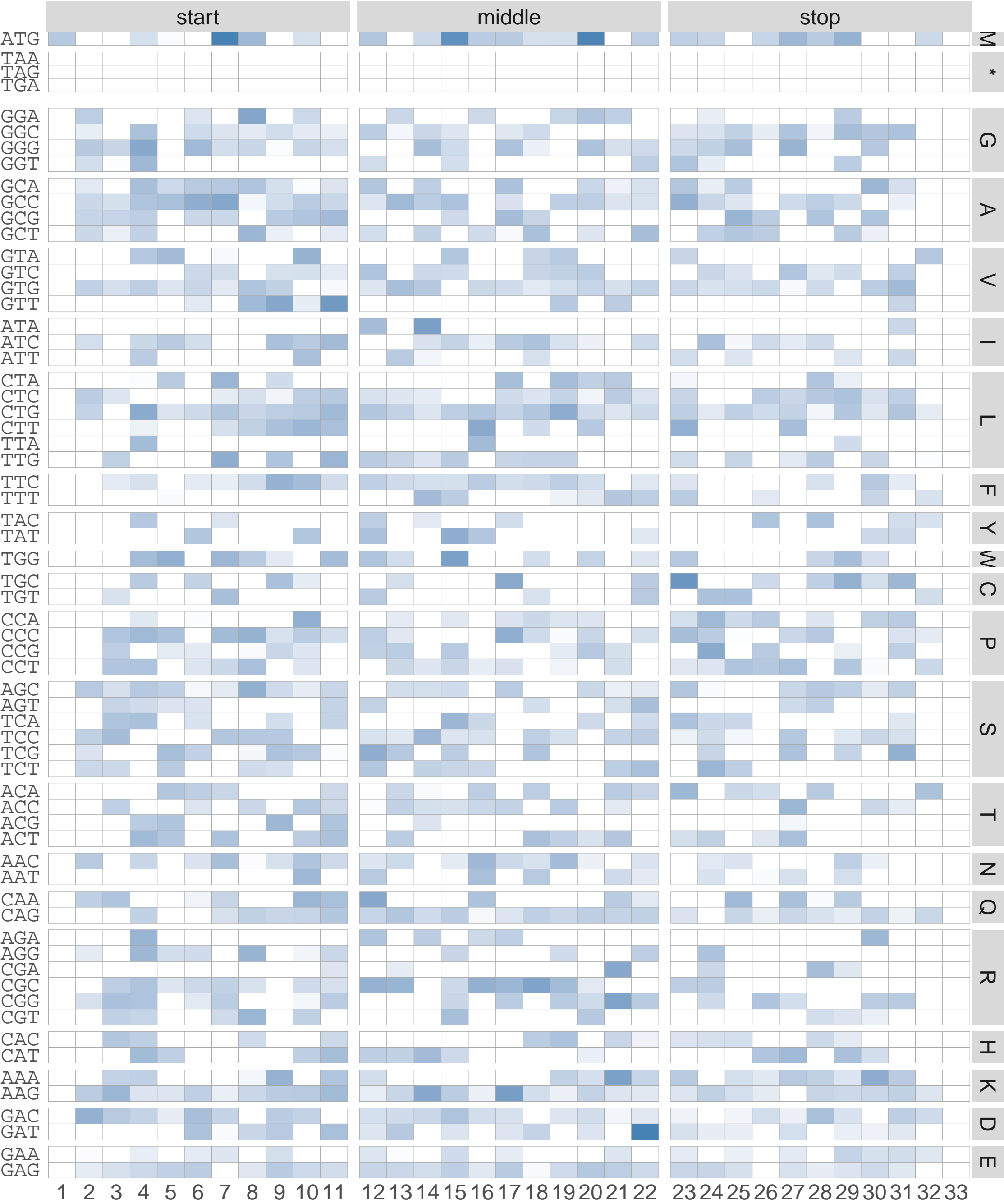sites per codon(scale: log2)