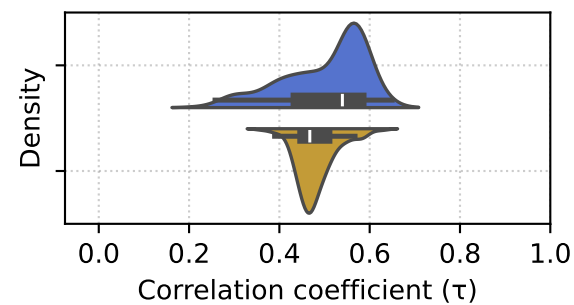
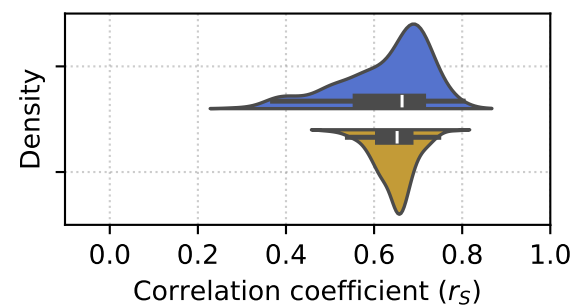
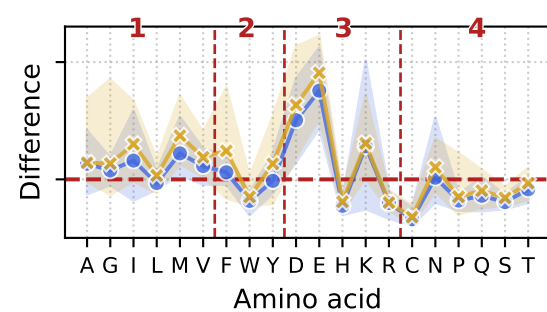
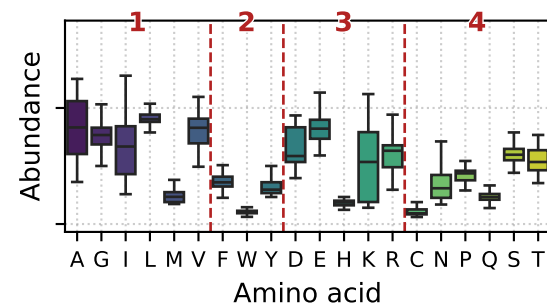
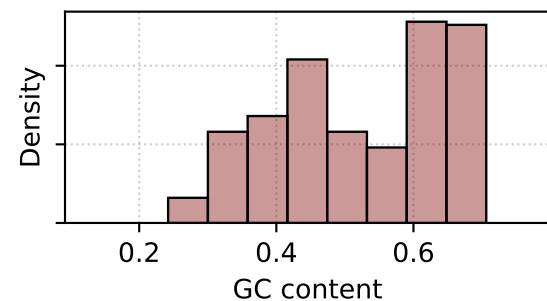
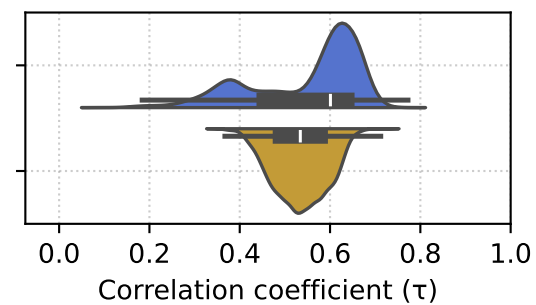
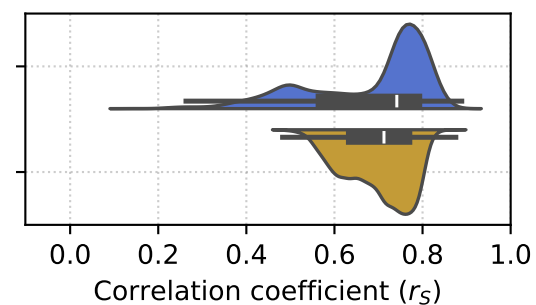
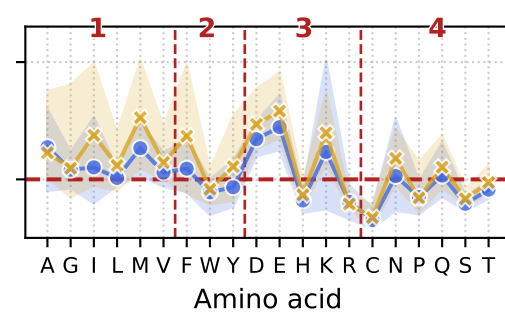
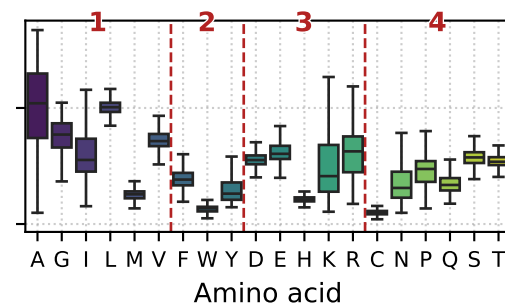
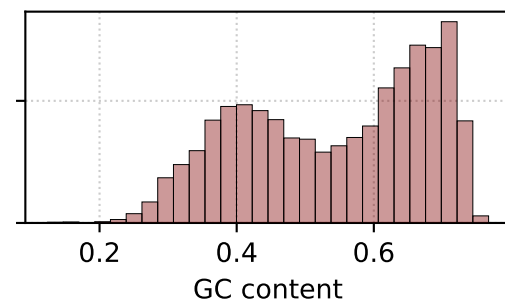


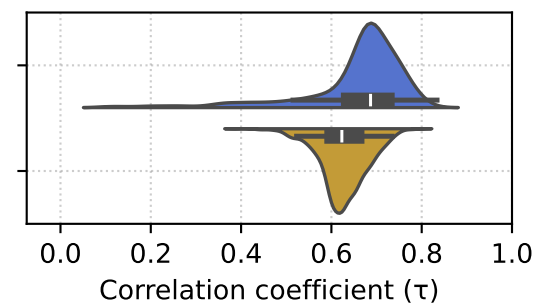
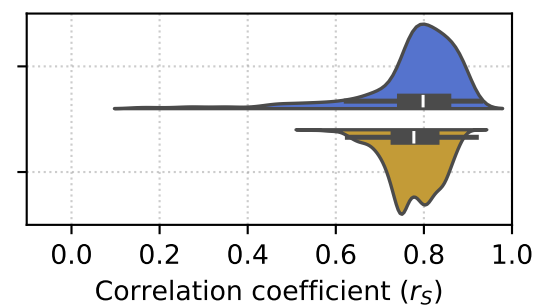
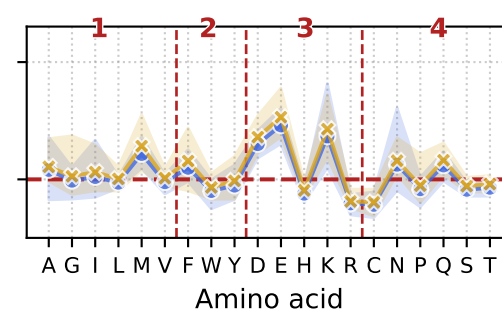
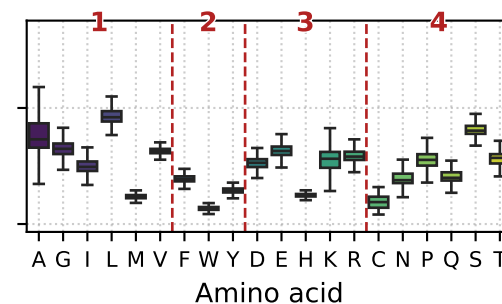
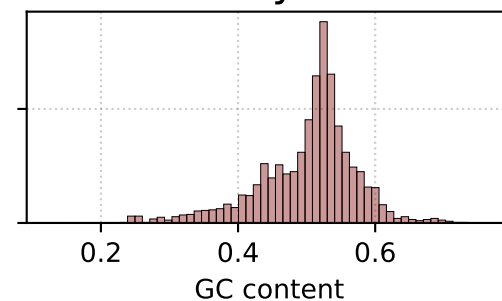
Archaea



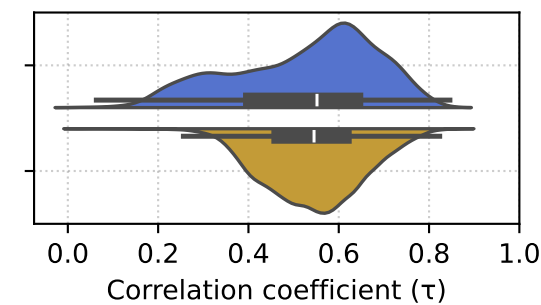
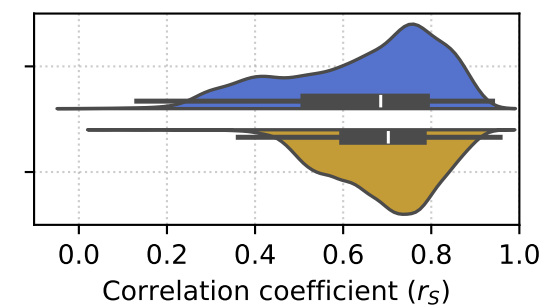
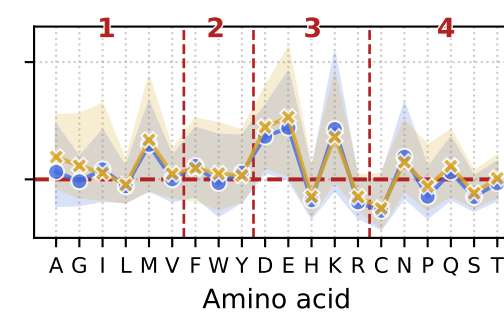
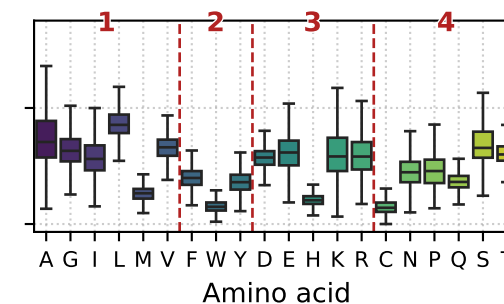
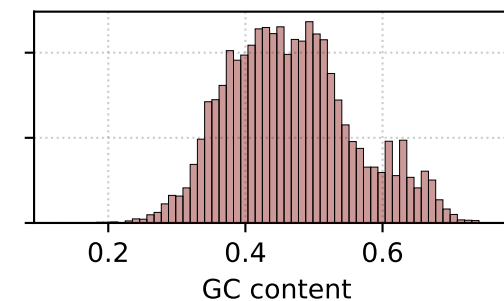
Bacteria



Eukaryotes



Viruses



Mean GC contents (%) of protein-coding genes per species

Mean amino acid abundances (%) per species

1 - Aliphatic
2 - Aromatic
3 - Charged
4 - Uncharged

Percentage differences (%) between empirical and expected abundances per species

1 - Aliphatic
2 - Aromatic
3 - Charged
4 - Uncharged

Codon number
Codon+GC

Spearman correlation coefficients per species

Codon number
Codon+GC

Kendall's Tau correlation coefficients per species

Codon number
Codon+GC