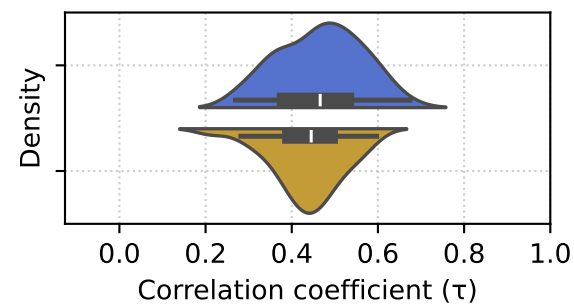
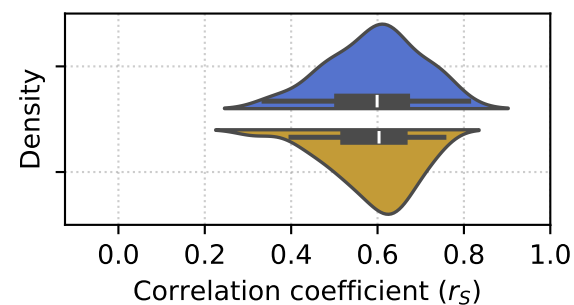
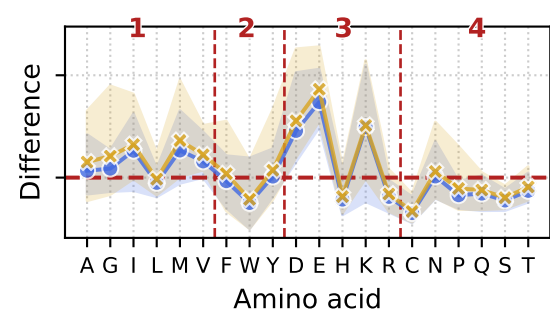
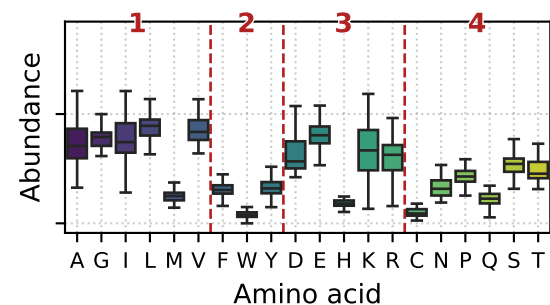
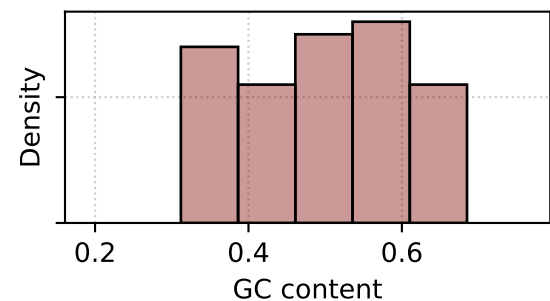
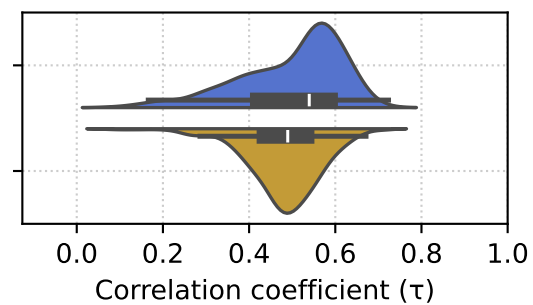
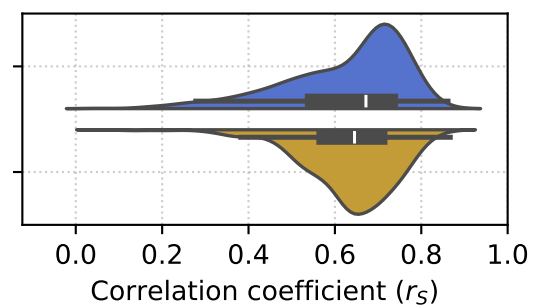
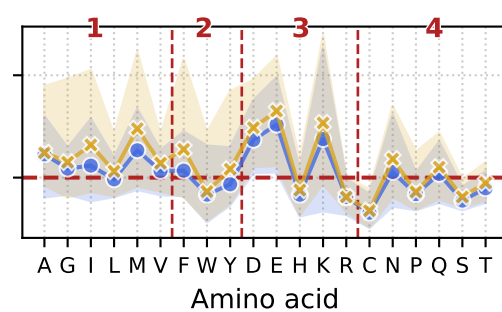
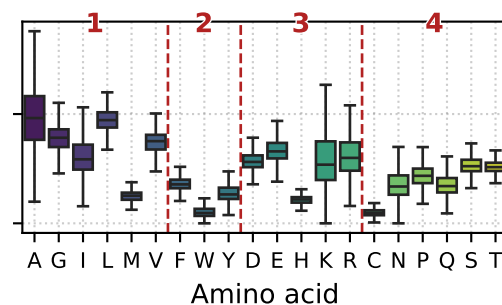
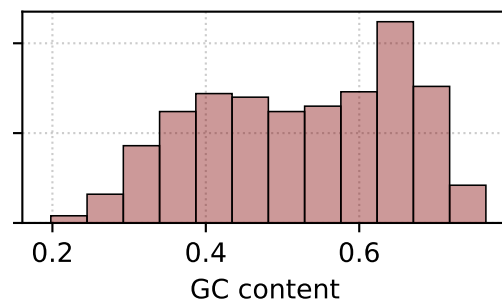


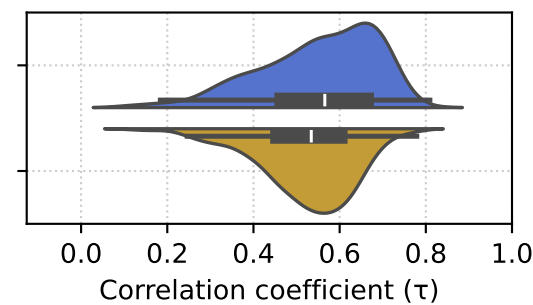
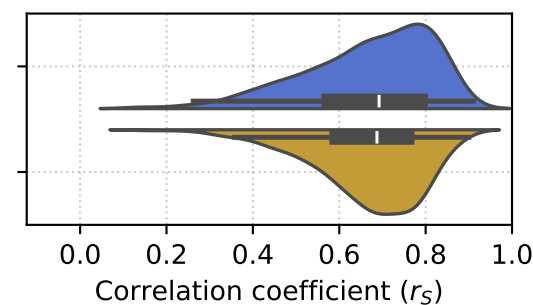
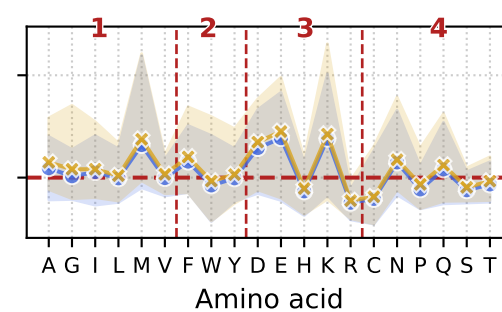
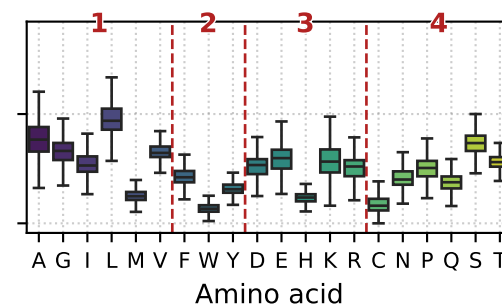
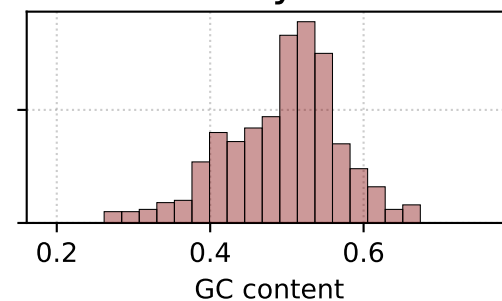
## Archaea



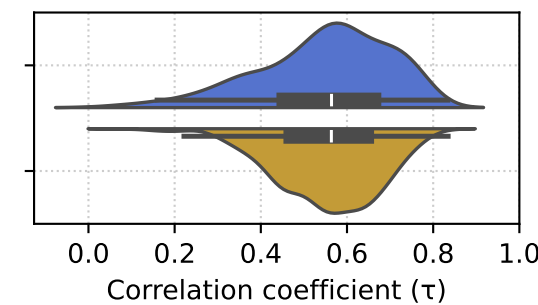
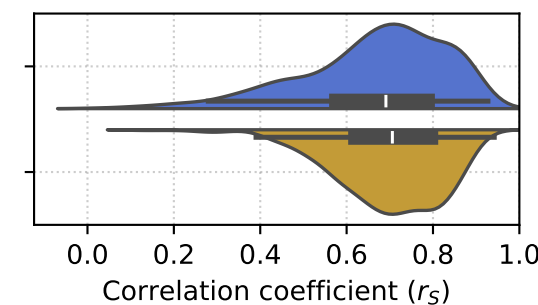
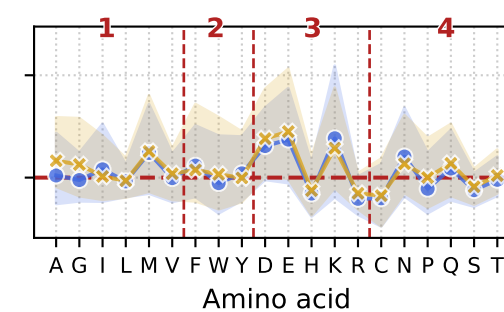
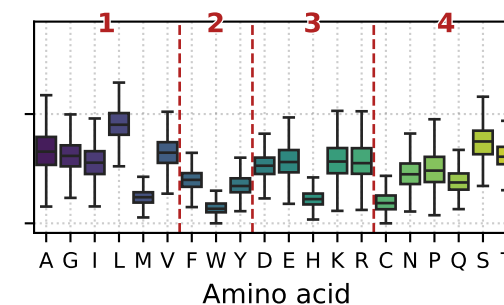
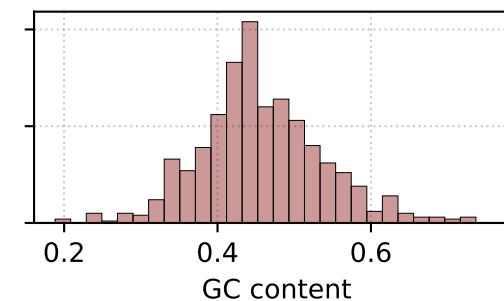
## Bacteria



## 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