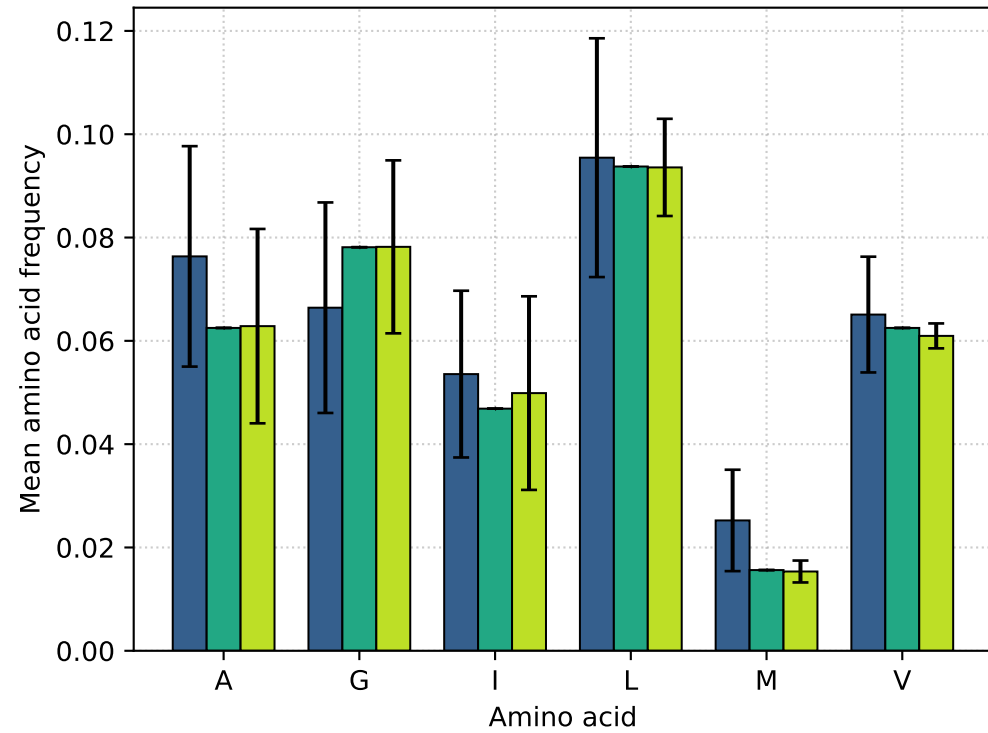


Eukaryota - Mean proteomic amino acid frequency for genetic code: Candidate division sr1 and gracilibacteria

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Pearson:
- Coefficient: 0.643526
- p-value: 0.024020

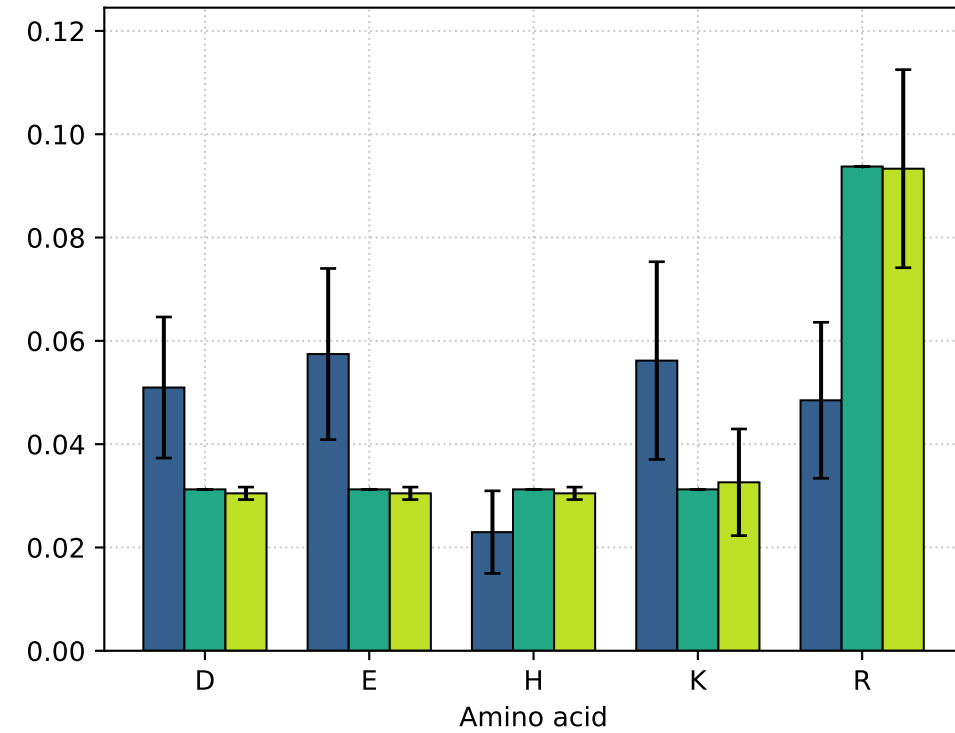
- Spearman:
- Coefficient: 0.665302
- p-value: 0.017168

Codon + GC correlation:

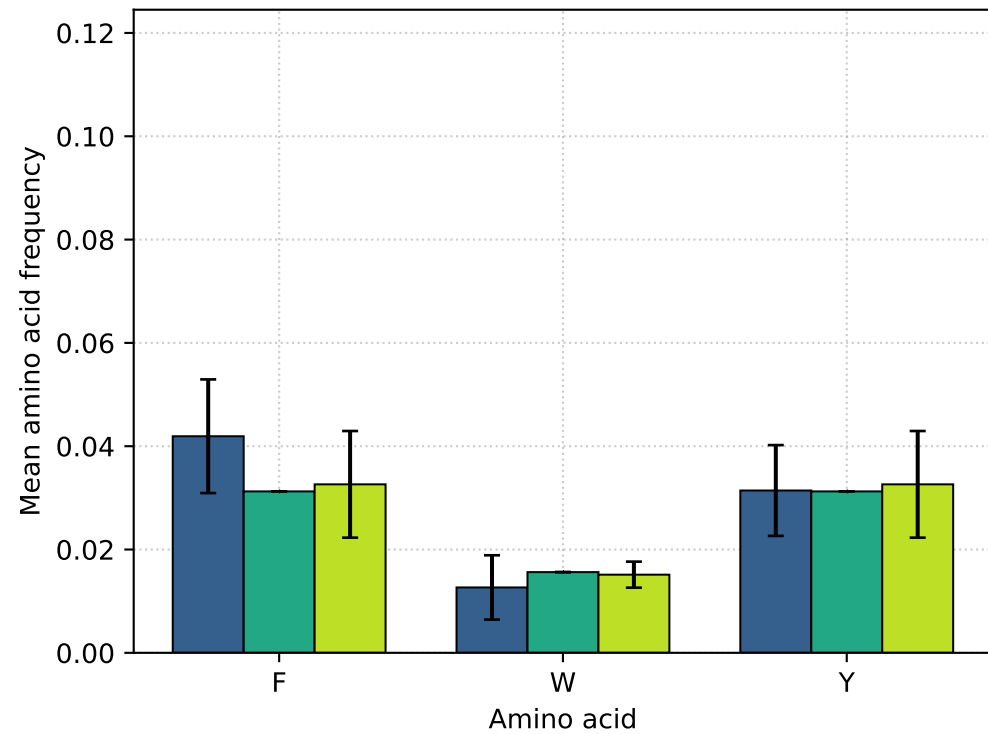
- Pearson:
- Coefficient: 0.672981
- p-value: 0.011907

- Spearman:
- Coefficient: 0.665586
- p-value: 0.010786

Charged amino acids



Aromatic amino acids



Uncharged amino acids

