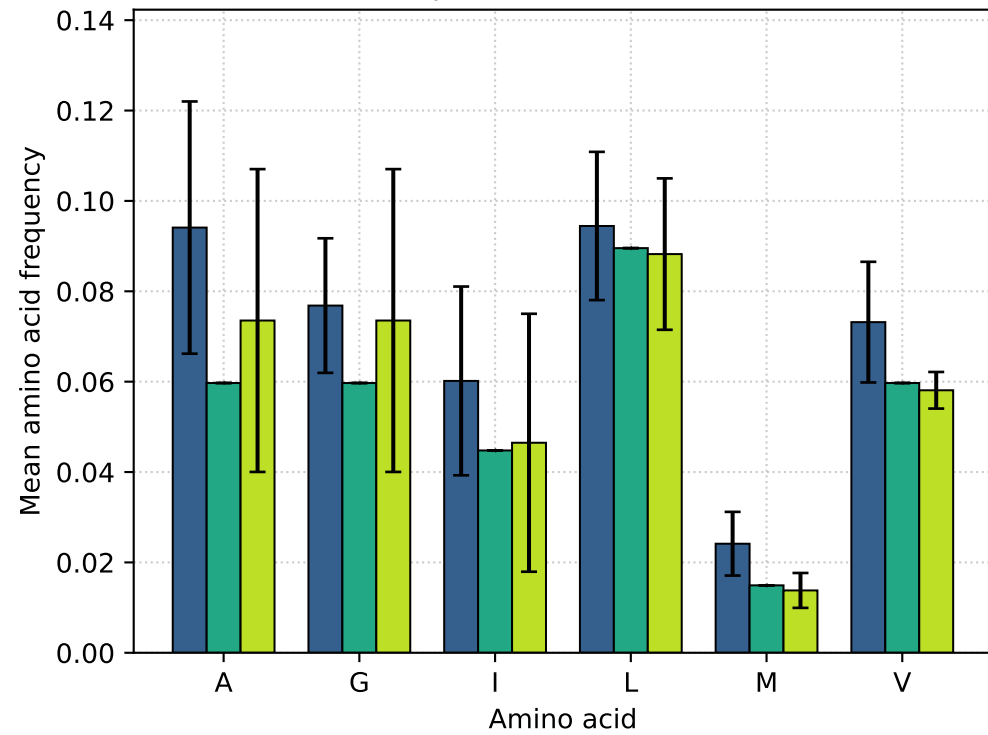


Bacteria - Mean proteomic amino acid frequency for genetic code: Condyllostoma nuclear

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Pearson:
- Coefficient: 0.509613
- p-value: 0.074180

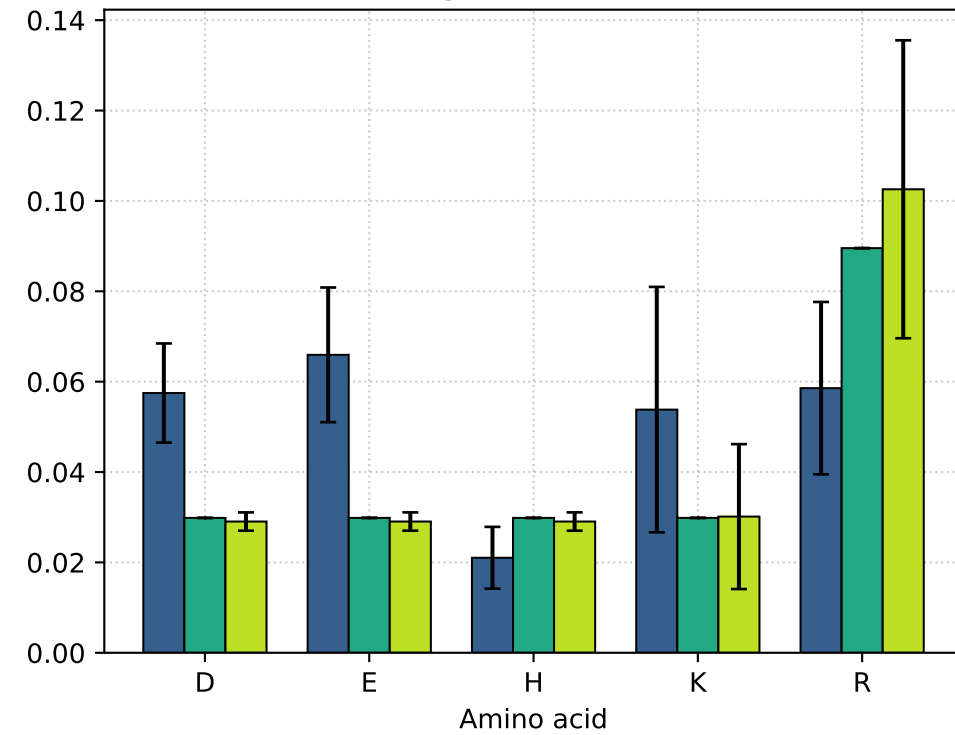
- Spearman:
- Coefficient: 0.544539
- p-value: 0.061374

Codon + GC correlation:

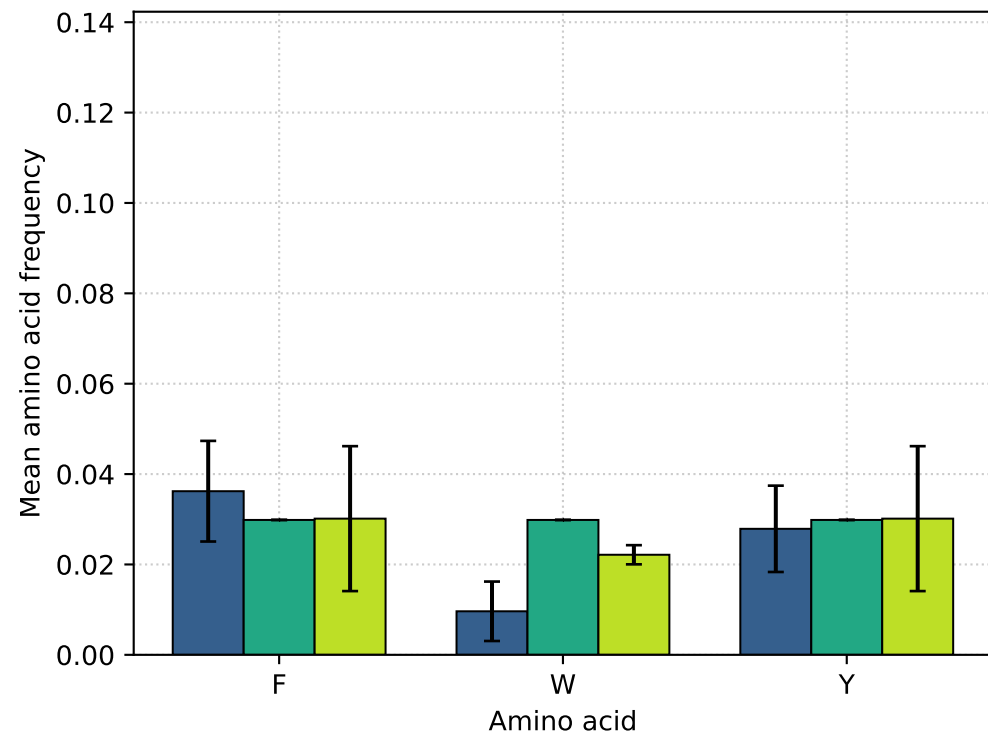
- Pearson:
- Coefficient: 0.610890
- p-value: 0.019780

- Spearman:
- Coefficient: 0.610950
- p-value: 0.020964

Charged amino acids



Aromatic amino acids



Uncharged amino acids

