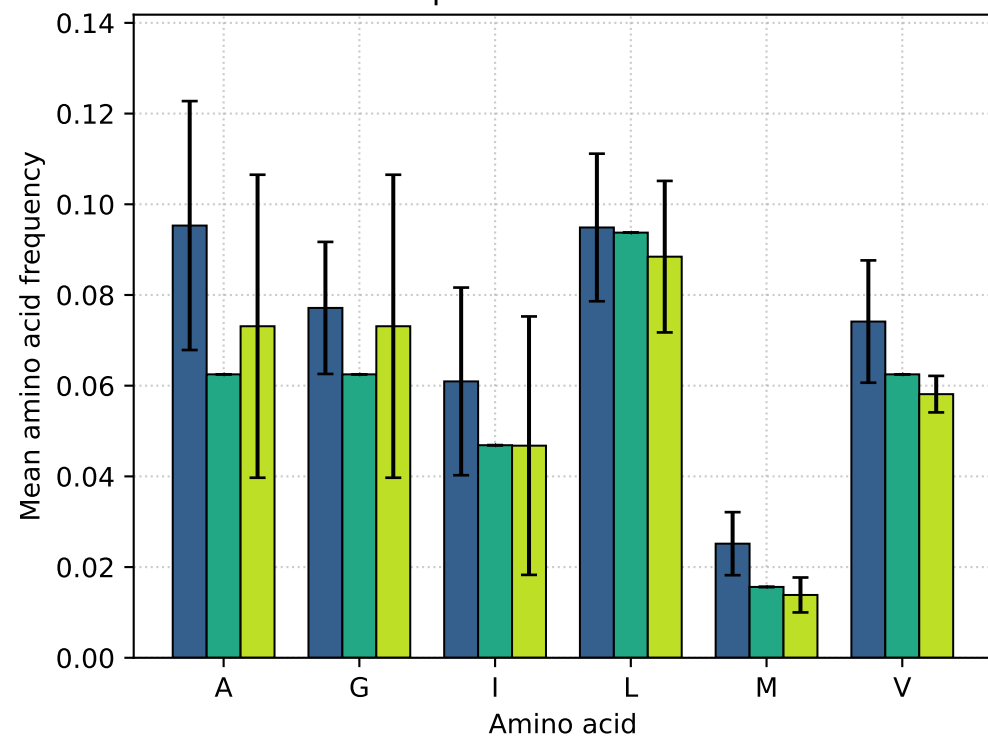


Bacteria - Mean proteome amino acid frequency for genetic code: Condyllostoma nuclear

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson: 0.51101
- Coefficient: 0.51101
- p-value: 7.25587e-02

Codon + GC correlation:

- Mean Pearson: 0.61027
- Coefficient: 0.61027
- p-value: 1.98768e-02

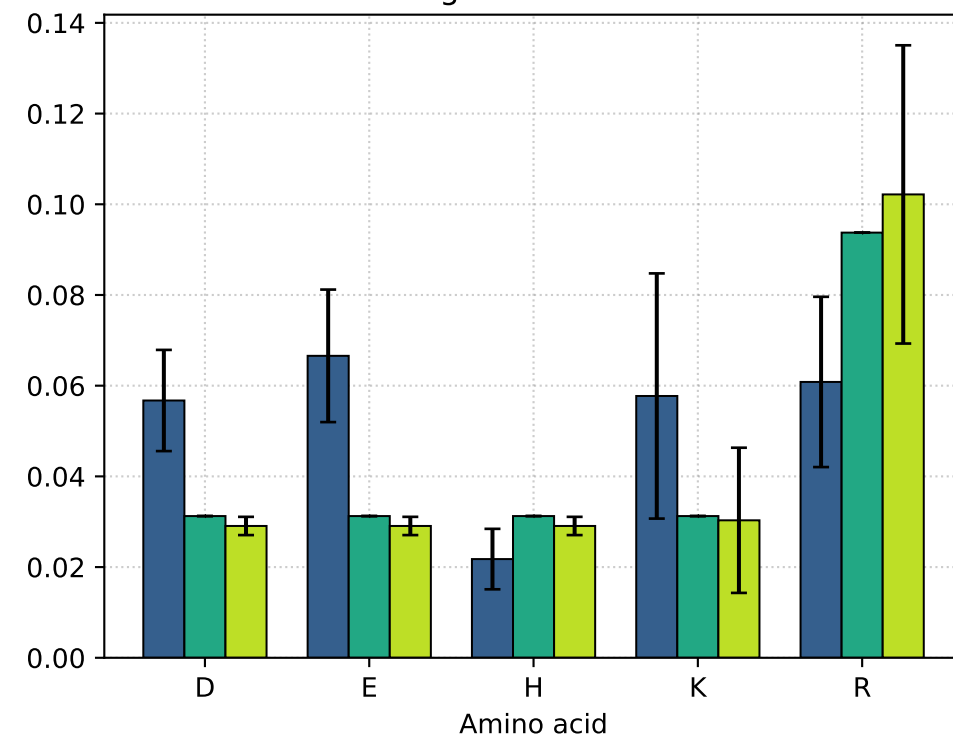
Mean Spearman:

- Coefficient: 0.54333
- p-value: 5.91741e-02

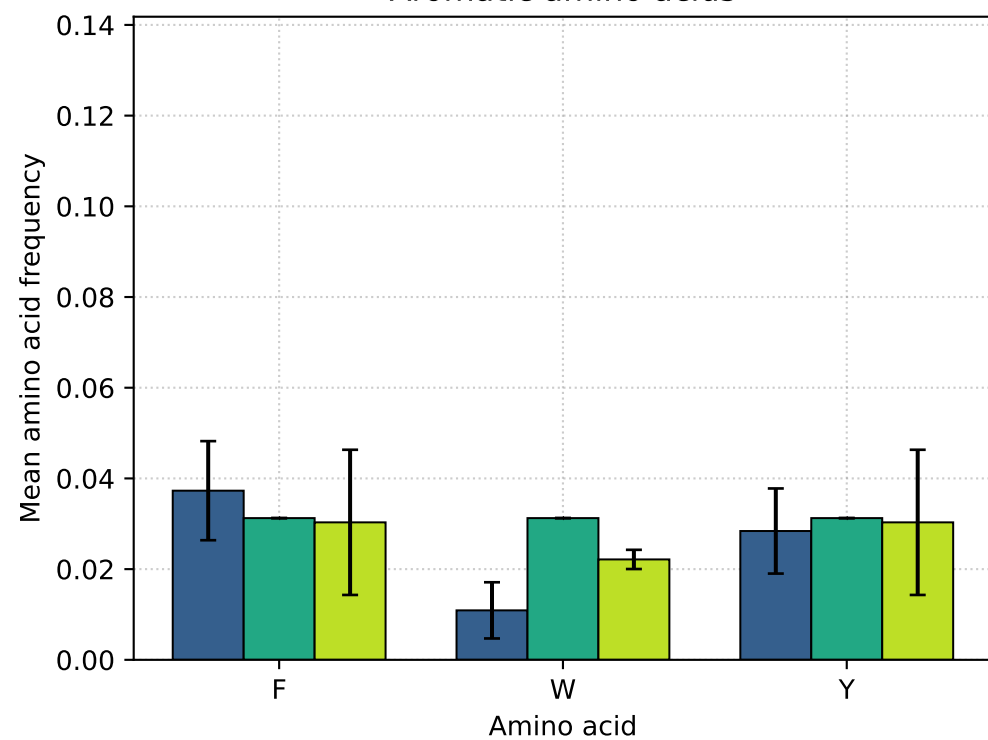
Mean Spearman:

- Coefficient: 0.60674
- p-value: 2.11956e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

